# **TC-Python Documentation**

Release 2025b

**Thermo-Calc Software AB** 

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### INSTALLATION GUIDE

This guide helps you to get a working TC-Python API installation.

There is a PDF guide included with your installation. In the Thermo-Calc menu, select  $Help \rightarrow Manuals$ Folder. Then double-click to open the Software Development Kits (SDKs) folder.

Note: A license is required to run TC-Python, you can read up details in *Configuring License*.

# 1.1 What type of installation should I choose?

There are two possibilities to install TC-Python:

- 1. Using the Python-interpreter bundled to Thermo-Calc: This interpreter has TC-Python preinstalled together with some popular Python-packages. This is the recommended option for new users to TC-Python, but it is limited to the preinstalled packages.
- 2. *Installing TC-Python into the Python-interpreter of your choice*: This is the recommended option for any more advanced usage and provides full flexibility.

# **1.2 Using the Python-interpreter bundled to Thermo-Calc**

Note: A Python-interpreter is bundled to Thermo-Calc beginning with version 2021a.

### 1.2.1 Limitations

The bundled Python 3 interpreter is containing the following major packages:

Package
colour-science
matplotlib
numpy
scikit-learn
scipy
pandas
pyvista
TC-Python
tmm

Please contact the Thermo-Calc support if you think that further packages might be useful in future releases.

**Note:** The following TC-Python examples are requiring additional packages that are not available in the bundled Python-interpreter, they can therefore not be run:

- pyex\_M\_01\_Input\_from\_file.py
- pyex\_M\_02\_Output\_to\_file.py

**Warning:** The Python-interpreter bundled to Thermo-Calc is also used for running the property models in Thermo-Calc. **Any changes to the interpreter packages can therefore break Thermo-Calc and should be avoided**. If the installation gets broken, it can be fixed by reinstalling Thermo-Calc after having removed it.

#### 1.2.2 Step 1: Install an IDE (Integrated Development Environment)

Any editor can be used to write the Python code, but an IDE is recommended, e.g. PyCharm. These instructions are based on the use of PyCharm.

Use of an IDE will give you access to code completion, which is of great help when you use the API as it will give you the available methods on the objects you are working with.

- 1. Navigate to the PyCharm website: https://www.jetbrains.com/pycharm/download.
- 2. Click to choose your OS and then click Download. You can use the Community version of PyCharm.
- 3. Follow the instructions. It is recommended you keep all the defaults.

**Note:** For Mac installations, you also need to set some environment variables as described below in *Setting Environment Variables*.

### 1.2.3 Step 2: Configure PyCharm to use the bundled Python-interpreter

Open PyCharm and configure the interpreter:

- 1. Go the menu **File** $\rightarrow$ **Settings**.
- 2. Navigate in the tree to Project: YourProjectName and choose Project Interpreter.
- 3. Click on the settings symbol close to the Project Interpreter dropdown menu and choose Add.
- 4. Now choose **System Interpreter** and add the bundled Thermo-Calc Python 3 interpreter. It is located in different places depending on the operating system:

Operating system	Path to the bundled Python-interpreter
Windows	C:\Program Files\Thermo-Calc\2025b\python\python.exe
Linux	<pre>/home/UserName/Thermo-Calc/2025b/python/bin/python3</pre>
MacOS	/Applications/Thermo-Calc-2025b.app/Contents/
	Resources/python/bin/python3

5. Select your added interpreter and confirm.

### 1.2.4 Step 3: Run a TC-Python Example

Now you are ready to start working with TC-Python.

It is recommended that you open one or more of the included examples to both check that the installation has worked and to start familiarizing yourself with the code.

#### 1.2.4.1 Open the TC-Python Project in PyCharm

When you first open the TC-Python project and examples, it can take a few moments for the Pycharm IDE to index before some of the options are available.

1. Open PyCharm and then choose **File**→**Open**. The first time you open the project you will need to navigate to the path of the TC-Python installation:

Operating	sys-	Path to the TC-Python folder
tem		
Windows		C:\Users\UserName\Documents\Thermo-Calc\2025b\SDK\
		TC-Python
Linux		/home/UserName/Thermo-Calc/2025b/SDK/TC-Python
MacOS		/Users/Shared/Thermo-Calc/2025b/SDK/TC-Python

- 2. Click on the Examples folder and then click OK.
- 3. From any subfolder:
  - Double-click to open an example file to examine the code.
  - Right-click an example and choose Run .

Note: If you are not following the recommended approach and create a *new* project (File $\rightarrow$ New **Project...**), you need to consider that by default the options to choose the interpreter are hidden within the

**Create Project** window. So click on **Project Interpreter: New Virtual Environment** and in most cases choose your *System Interpreter* containing the Python bundled to Thermo-Calc instead of the default *New Virtual Environment*.

# 1.3 Installing TC-Python into the Python-interpreter of your choice

#### 1.3.1 Step 1: Install a Python Distribution

If you already have a Python distribution installation, version 3.8 or higher, skip this step.

These instructions are based on using the Anaconda platform for the Python distribution. Install version 3.8 or higher to be able to work with TC-Python, although it is recommended that you use the most recent version.

Note: TC-Python requires Python 3.8 or newer.

#### 1.3.1.1 Install Anaconda

- 1. Navigate to the Anaconda website: https://www.anaconda.com/download/.
- 2. Click to choose your OS (operating system) and then click **Download**. Follow the instructions. It is recommended you keep all the defaults.

#### **1.3.2 Step 2: Install Thermo-Calc and the TC-Python SDK**

Note: TC-Python is available starting with Thermo-Calc version 2018a.

- 1. Install Thermo-Calc
- 2. When the installation is complete, open the TC-Python folder that includes the file:\*.*whl* file needed for the next step. There is also an file:*Examples* folder with Python files you can use in the IDE to understand and work with TC-Python.

#### 1.3.3 Step 3: Install TC-Python

On Windows, it is recommended that you use the Python distribution prompt (i.e. Anaconda, ...), especially if you have other Python installations. Do not use Virtual Environments unless you have a good reason for that.

- 1. Open the command line. For example, in Anaconda on a Windows OS, go to **Start→Anaconda→Anaconda Prompt**.
- 2. At the command line, enter the following. Make sure there are no spaces at the end of the string or in the folder name or it will not run:

**Tip:** Note that on Linux depending on the interpreter usually *pip3* is used.

Operating	sys-	Path to the TC-Python folder
tem		
Windows		C:\Users\UserName\Documents\Thermo-Calc\2025b\SDK\
		TC-Python
Linux		/home/UserName/Thermo-Calc/2025b/SDK/TC-Python
MacOS		/Users/Shared/Thermo-Calc/2025b/SDK/TC-Python

3. Press <Enter>. When the process is completed, there is a confirmation that TC-Python is installed.

**Note:** If your computer is located behind a proxy-server, the default *pip*-command will fail with a network connection error. In that case you need to install the dependencies of TC-Python in a special configuration:

pip install -proxy user:password@proxy\_ip:port py4j jproperties

See "pip install" fails with "Failed to establish a new network connection" or similar for detailed information.

#### 1.3.4 Step 4: Install an IDE (Integrated Development Environment)

Any editor can be used to write the Python code, but an IDE is recommended, e.g. PyCharm. These instructions are based on the use of PyCharm.

Use of an IDE will give you access to code completion, which is of great help when you use the API as it will give you the available methods on the objects you are working with.

- 1. Navigate to the PyCharm website: https://www.jetbrains.com/pycharm/download.
- 2. Click to choose your OS and then click **Download**. You can use the **Community** version of PyCharm.
- 3. Follow the instructions. It is recommended you keep all the defaults.

**Note:** For Mac installations, you also need to set some environment variables as described below in *Setting Environment Variables*.

#### 1.3.5 Step 5: Open the IDE and Run a TC-Python Example

After you complete all the software installations, you are ready to open the IDE to start working with TC-Python.

It is recommended that you open one or more of the included examples to both check that the installation has worked and to start familiarizing yourself with the code.

#### 1.3.5.1 Open the TC-Python Project in PyCharm

When you first open the TC-Python project and examples, it can take a few moments for the Pycharm IDE to index before some of the options are available.

1. Open PyCharm and then choose **File**→**Open**. The first time you open the project you will need to navigate to the path of the TC-Python installation.

Operating	sys-	Path to the TC-Python folder
tem		
Windows		C:\Users\UserName\Documents\Thermo-Calc\2025b\SDK\
		TC-Python
Linux		/home/UserName/Thermo-Calc/2025b/SDK/TC-Python
MacOS		/Users/Shared/Thermo-Calc/2025b/SDK/TC-Python

- 2. Click on the Examples folder and then click **OK**.
- 3. From any subfolder:
  - Double-click to open an example file to examine the code.
  - Right-click an example and choose Run .

#### 1.3.5.2 Fixing potential issues with the environment

In most cases you should run TC-Python within your **global** Python 3 interpreter and not use Virtual Environments unless you have a good reason to do so. A common problem on first usage of TC-Python is the error message "**No module named tc\_python**". You can resolve this and other problems with the interpreter settings as follows:

- 1. Go the menu **File** $\rightarrow$ **Settings**.
- 2. Navigate in the tree to **Project: YourProjectName** and choose **Project Interpreter**.
- 3. Click on the settings symbol close to the **Project Interpreter** dropdown menu and choose Add.
- 4. Now choose System Interpreter and add your existing Python 3 interpreter.
- 5. Select your added interpreter and confirm.

**Note:** If you are not following the recommended approach and create a *new* project (File $\rightarrow$ New **Project...**), you need to consider that by default the options to choose the interpreter are hidden within the **Create Project** window. So click on **Project Interpreter:** New Virtual Environment and in most cases choose your *System Interpreter* instead of the default *New Virtual Environment*.

**Note:** If you really need to use a Virtual Environment, please consider the hints given in the *Python Virtual Environments* chapter.

# 1.4 Optional packages

Plotting results of Additive Manufacturing simulations requires the package *pyvista*. It is automatically installed if using the option [*am-plotting*] during the installation:

The bundled interpreter contains *pyvista* by default.

# 1.5 Updating to a newer version

When updating to a newer version of Thermo-Calc, **you need to also install the latest version of TC-Python**. This is not necessary if are using the bundled Python-interpreter that has it automatically installed. It is not sufficient to run the installer of Thermo-Calc:

pip install <path to the TC-Python folder>/TC\_Python-<version>-py3-none-any.whl

**Tip:** Note that on Linux depending on the interpreter usually *pip3* is used.

In case of problems you may wish to uninstall the previous version of TC-Python in advance:

```
pip uninstall TC-Python
pip install rone-any.whl
```

However, that should normally not be required.

You can check the currently installed version of TC-Python by running:

pip show TC-Python

# **CONFIGURING LICENSE**

TC-Python requires a license to run, you can obtain a license from the Thermo-Calc support @thermocalc.com.

If you have no valid license, an exception will be thrown that states: *NO LICENSE: No activated license for TC-Python*. Depending on the type of license, different steps are required to configure TC-Python to use it.

# 2.1 Environment variables

Some environment variables need to be set. On Windows this is normally done during the installation of Thermo-Calc, but on Linux and macOS this needs to be performed for each used IDE or terminal. Details are described in the chapter *Setting Environment Variables*.

# 2.2 License activation

#### Only required for licenses with user credentials.

You need to activate the TC-Python license once on your computer. *This license differs from that for the Thermo-Calc application and thus it is not sufficient to activate only the Thermo-Calc GUI.* 

The activation is done with the methods in the class tc\_python.license.LicenseManager. Depending on the type of license various methods are available, for example for online and offline activation.

It is recommended to simply run the example Miscellaneous/pyex\_M\_04\_license.py to perform the activation and deactivation. By default an online activation is performed, but you can choose another method by activating another method among those being commented out at the end of the file.

Note: You need to deactivate the license if you want to use it on another computer.

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# SETTING ENVIRONMENT VARIABLES

In order to use TC-Python on macOS and Linux you need to set some environment variables.

TC25B\_HOME=/Applications/Thermo-Calc-2025b.app/Contents/Resources

If you use a license server:

LSHOST=<name-of-the-license-server>

If you have a node-locked license:

LSHOST=NO-NET LSERVRC=/Users/Shared/Thermo-Calc/lservrc

If you have a license based on user credentials:

TC\_LICENSE\_SPRING=y

If running in the terminal they can be set using *export LSHOST=NO-NET*, etc.

In PyCharm, you can add environment variables in the configurations.

Select  $Run \rightarrow Edit$  Configurations to open the Run/Debug Configurations window. Choose Templates and then Python. Enter the environment variable(s) by clicking the button to the right of the Environment Variables text field. Now the environment variables(s) will be set for each new configuration by default.

**Note:** Existing configurations need to be removed and recreated to obtain the environment variables in them.

The same way for configuring the environment variables can be used on other operating systems as if necessary.

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### **ARCHITECTURE OVERVIEW**

TC-Python contains classes of these types:

- TCPython this is where you start with general settings.
- SystemBuilder and System where you choose database and elements etc.
- Calculation where you choose and configure the calculation.
- **Result** where you get the results from a calculation you have run.

### 4.1 TCPython

This is the starting point for all TC-Python usage.

You can think of this as the start of a "wizard".

You use it to select databases and elements. That will take you to the next step in the wizard, where you configure the system.

**Example:** 

```
from tc_python import *
with TCPython() as start:
    start.select_database_and_elements(...
    # e.t.c
# after with clause
# or like this
with TCPython():
    SetUp().select_database_and_elements(...
    # e.t.c
# after with clause
```

**Tip:** If you use TC-Python from Jupyter Lab / Notebook, you should use TC-Python slightly different to be able to use multiple cells. See *Using TC-Python within a Jupyter Notebook or the Python console* for details.

Note: When your python script runs a row like this:

with TCPython() as start:

a process running a calculation server starts. Your code, via TC-Python, uses socket communication to send and receive messages to and from that server.

When your Python script has run as far as this row:

# after with clause

the calculation server automatically shuts down, and all temporary files are deleted. It is important to ensure that this happens by structuring your Python code using a with() clause as in the above example.

**Note:** To re-use results from previous calculations, set a folder where TC-Python saves results, and looks for previous results.

This is done with the function set\_cache\_folder().

```
from tc_python import *
with TCPython() as start:
    start.set_cache_folder("cache")
```

This folder can be a network folder and shared by many users. If a previous TC-Python calculation has run with the same cache\_folder and EXACTLY the same system and calculation settings, the calculation is not re-run. Instead the result is automatically loaded from disk.

It is also possible to explicitly save and load results.

```
from tc_python import *
with TCPython() as start:
    #... diffusion calculation (could be any calculation type)
    calculation_result.save_to_disk('path to folder')
    #...
    loaded_result = start.load_result_from_disk().diffusion('path to folder')
```

### 4.2 SystemBuilder and System

A SystemBuilder is returned when you have selected your database and elements in TCPython.

The **SystemBuilder** lets you further specify your system, for example the phases that should be part of your system.

**Example:** 

```
from tc_python import *
with TCPython() as start:
    start.select_database_and_elements("ALDEMO", ["Al", "Sc"])
    # e.t.c
```

When all configuration is done, you call get\_system() which returns an instance of a System class. The System class is fixed and cannot be changed. If you later want to change the database, elements or something else, change the SystemBuilder and call get\_system() again, or create a new SystemBuilder and call get\_system().

From the **System** you can create one or more calculations, which is the next step in the "wizard".

Note: You can use the same System object to create several calculations.

# 4.3 Calculation

The best way to see how a calculation can be used is in the TC-Python examples included with the Thermo-Calc installation.

Some calculations have many settings. Default values are used where it is applicable, and are overridden if you specify something different.

When you have configured your calculation you call calculate() to start the actual calculation. That returns a **Result**, which is the next step.

### 4.3.1 Single equilibrium calculations

In single equilibrium calculations you need to specify the correct number of conditions, depending on how many elements your **System** contains.

You do that by calling set\_condition().

An important difference from other calculations is that single equilibrium calculations have two functions to get result values.

The calculate() method, which gives a **SingleEquilibriumTempResult**, is used to get actual values. This result is "temporary", meaning that if you run other calculations or rerun the current one, the resulting object no longer gives values corresponding to the first calculation.

This is different from how other calculations work. If you want a **Result** that you can use *after* running other calculations, you need to call calculate\_with\_state(), which returns a **SingleEquilibriumResult**.

**Note:** calculate() is the recommended function and works in almost all situations. Also it has *much* better performance than calculate\_with\_state().

```
from tc_python import *
with TCPython() as start:
    gibbs_energy = (
        start.
        select_database_and_elements("FEDEMO", ["Fe", "Cr", "C"]).
        get_system().
        with_single_equilibrium_calculation().
            set_condition(ThermodynamicQuantity.temperature(), 2000.0).
            set_condition(ThermodynamicQuantity.mole_fraction_of_a_component("Cr"),_L
            set_condition(ThermodynamicQuantity.mole_fraction_of_a_component("C"), 0.
            set_condition(ThermodynamicQuantity.mole_fraction_of_a_component("C"), 0.
            set_value_of("G")
            )
```

### 4.3.2 Batch equilibrium calculations

Batch equilibrium calculations are used when you want to do many single equilibrium calculations and it is known from the beginning which result values are required from the equilibrium. This is a vectorized type of calculation that can reduce the overhead from Python and TC-Python similar to the approach used in *numpy*-functions for example.

**Tip:** The performance of batch equilibrium calculations can be significantly better than looping and using single equilibrium calculations **if the actual Thermo-Calc calculation is fast**. There is little advantage if the Thermo-Calc equilibrium calculations take a long time (typically for large systems and databases).

```
from tc_python import *
with TCPython() as start:
   calculation = (
        start
            .set_cache_folder(os.path.basename(__file__) + "_cache")
            .select_database_and_elements("NIDEMO", ["Ni", "Al", "Cr"])
            .get_system()
            .with_batch_equilibrium_calculation()
            .set_condition("T", 800.1)
            .set_condition("X(Al)", 1E-2)
            .set_condition("X(Cr)", 1E-2)
            .disable_global_minimization()
   )
   list_of_x_Al = [1, 2, 3, 4, 5, 6, 7, 8, 9, 10]
   list_of_x_Cr = [3, 5, 7, 9, 11, 13, 15]
   lists_of_conditions = []
    for x_Al in list_of_x_Al:
        for x_Cr in list_of_x_Cr:
            lists_of_conditions.append([
                ("X(Al)", x_Al / 100),
                ("X(Cr)", x_Cr / 100)])
    calculation.set_conditions_for_equilibria(lists_of_conditions)
   results = calculation.calculate(["BM", "VM"])
   masses = results.get_values_of("BM")
   volumes = results.get_values_of('VM')
print(masses)
print(volumes)
```

### 4.3.3 Precipitation calculations

All that can be configured in the *Precipitation Calculator* in Graphical Mode can also be done here in this calculation. However, you must at least enter a matrix phase, a precipitate phase, temperature, simulation time and compositions.

**Example:** 

### 4.3.4 Scheil calculations

All Scheil calculations available in Graphical Mode or Console Mode can also be done here in this calculation. The minimum you need to specify are the elements and compositions. Everything else is set to a default value.

### 4.3.5 Property diagram calculations

For the property diagram (step) calculation, everything that you can configure in the *Equilibrium Calculator* when choosing *One axis* in Graphical Mode can also be configured in this calculation. In Console Mode the property diagram is created using the Step command. The minimum you need to specify are elements, conditions and the calculation axis. Everything else is set to default values, if you do not specify otherwise.

**Example:** 

```
from tc_python import *
with TCPython() as start:
    property_diagram = (
        start.
            select_database_and_elements("FEDEMO", ["Fe", "C"]).
            get_system().
            with_property_diagram_calculation().
                with_axis(CalculationAxis(ThermodynamicQuantity.temperature()).
                    set_min(500).
                    set_max(3000)).
                set_condition(ThermodynamicQuantity.mole_fraction_of_a_component("C"), 0.
→01).
                calculate().
                get_values_grouped_by_stable_phases_of(ThermodynamicQuantity.
\rightarrow temperature(),
                                                         ThermodynamicQuantity.volume_
→ fraction_of_a_phase("ALL"))
        )
```

### 4.3.6 Phase diagram calculations

For the phase diagram (map) calculation, everything that you can configure in the *Equilibrium Calculator* when choosing *Phase diagram* in Graphical Mode can also be configured in this calculation. In Console Mode the phase diagram is created using the Map command. The minimum you need to specify are elements, conditions and two calculation axes. Everything else is set to default values, if you do not specify otherwise.

**Example:** 

### 4.3.7 Diffusion calculations

For diffusion calculations, everything that you can configure in the *Diffusion Calculator* can also be configured in this calculation. The minimum you need to specify are elements, temperature, simulation time, a region with a grid and width, a phase and an initial composition.

**Example:** 

```
from tc_python import *
with TCPython() as start:
    diffusion_result = (
        start.
            select_thermodynamic_and_kinetic_databases_with_elements("FEDEMO", "MFEDEMO",
\rightarrow ["Fe", "Ni"]).
            get_system().
            with_isothermal_diffusion_calculation().
                set_temperature(1400.0).
                set_simulation_time(108000.0).
                add_region(Region("Austenite").
                    set_width(1E-4).
                    with_grid(CalculatedGrid.linear().set_no_of_points(50)).
                    with_composition_profile(CompositionProfile().
                         add("Ni", ElementProfile.linear(10.0, 50.0))
                    ).
                add_phase("FCC_A1")).
        calculate())
    distance, ni_fraction = diffusion_result.get_mass_fraction_of_component_at_time("Ni",
→ 108000.0)
```

### 4.3.8 Property Model calculations

For Property Model calculations, everything that you can configure in the *Property Model Calculator* in Graphical Mode can also be configured in this calculation. The minimum you need to specify are elements, composition and which Property Model you want to use.

**Example:** 

```
from tc_python import *
with TCPython() as start:
```

```
print("Available Property Models: {}".format(start.get_property_models()))
property_model = (
    start.
        select_database_and_elements("FEDEMO", ["Fe", "C"]).
        get_system().
        with_property_model_calculation("Driving force").
        set_composition("C", 1.0).
        set_argument("precipitate", "GRAPHITE"))
print("Available arguments: {}".format(property_model.get_arguments()))
result = property_model.calculate()
print("Available result quantities: {}".format(result.get_result_quantities()))
driving_force = result.get_value_of("normalizedDrivingForce")
```

### 4.3.9 Material to Material calculations

Material to Material calculations are generally regular single equilibrium, property diagram or phase diagram calculations but they are specialised to handle the mixture of two materials A and B. Everything that you can configure in the *Material to Material Calculator* in Graphical Mode can also be configured in this calculation. The minimum required configuration is shown below for a *Property diagram calculation for varying amount of material B*. The other calculators (single fraction of material B and phase diagram calculations) are configured in a similar way.

#### Example:

```
from tc_python import *
with TCPython() as start:
    material_to_material_property_diagram = (
        start.
            select_database_and_elements("FEDEMO", ["Fe", "Cr", "Ni", "C"]).
            get_system().
            with_material_to_material().
            with_property_diagram_calculation().
            set_material_a({"Cr": 10.0, "Ni": 15.0}, "Fe").
            set_material_b({"Cr": 15.0, "Ni": 10.0}, "Fe").
            set_activities({"C": 0.1}).
            with_constant_condition(ConstantCondition.temperature(800 + 273.15)).
            with_axis(MaterialToMaterialCalculationAxis.fraction_of_material_b(from_
\rightarrow fraction=0.0,
                                                                                    to_
\rightarrow fraction=1.0,
                                                                                    start
\rightarrow fraction=0.5))
    )
    result = material_to_material_property_diagram.calculate()
    data = result.get_values_grouped_by_quantity_of(MATERIAL_B_FRACTION,
                                                       ThermodynamicQuantity.volume_

→fraction_of_a_phase(ALL_PHASES))
```

```
for group in data.values():
    fractions_of_b = group.x
    volume_fractions_of_phase = group.y
    phase_name = group.label
```

#### 4.3.10 Process Metallurgy calculations

Process Metallurgy calculations are specialized to support the convenient handling of component-based additions (i.e., slag compositions such as 50% Al2O3 - 30% CaO - 20% SiO2), provide tailor-made result quantities, a framework for developing kinetic process simulations, and more useful features.

There are two distinct types of calculations:

- tc\_python.process\_metallurgy.equilibrium.EquilibriumCalculation: isothermal and adiabatic equilibrium calculations
- tc\_python.process\_metallurgy.process.ProcessSimulationCalculation: a kinetic process simulation framework, based on an *Effective Equilibrium Reaction Zone* (EERZ) approach

#### Equilibrium calculation example:

Equilibrium calculations are useful in a large range of situations when considering the kinetics of a process is unnecessary.

```
from tc_python import *
with TCPython() as session:
    metal = EquilibriumAddition({"Fe": None, "C": 4.5, "Si": 1.0}, 100e3,...
\rightarrow temperature=1650 + 273.15)
    slag = EquilibriumAddition({"Ca0": 75, "Al203": 25}, 3e3, temperature=1600 + 273.15)
    gas = EquilibriumGasAddition({"02": 100}, 1000, amount_unit=GasAmountUnit.NORM_CUBIC_
\rightarrow METER)
    calc = session.with_metallurgy().with_adiabatic_equilibrium_
→calculation(ProcessDatabase.OXDEMO)
    (calc
     .add_addition(metal)
     .add_addition(slag)
     .add_addition(gas))
    result = calc.calculate()
    print(f"Stable phases: {result.get_stable_phases()}, temperature: {result.get_
\rightarrow temperature() } K")
```

#### **Process simulation example:**

TC-Python is providing a *framework* for modelling in principle any process in metallurgy, especially steel-making. It is up to the user to actually develop a concrete model for the process in question. The framework is in the current release limited to one reaction zone connecting two bulk zones. These bulk zones are typically the steel melt and the top slag, but not limited to that. The framework in its current version has proven to be useful to model industrial ladle furnaces, AOD- and VOD-converters and more. Process features such as heating and cooling, heat transfer between the bulk zones, inclusion formation and their flotation, etc., can be modelled.

This is a very simplified minimal but complete model mimicking a BOF process:

```
from tc_python import *
with TCPython() as session:
    calc = (session.with_metallurgy()
             .with_adiabatic_process_calculation(ProcessDatabase.OXDEMO)
            .set_end_time(15 * 60))
    steel_zone = MetalBulkZone(density=7800)
    slag_zone = SlagBulkZone(density=4500)
    steel_zone.add_addition(SingleTimeAddition({"Fe": None, "C": 4.5, "Si": 1.0}, 120e3,
                                                  temperature=1600 + 273.15), time=0)
    slag_zone.add_addition(SingleTimeAddition({"Ca0": 75, "Si02": 25}, 1.2e3,
                                                temperature = 1500 + 273.15,
                                                 composition_unit=CompositionUnit.MOLE_
\rightarrow PERCENT), time=\emptyset)
    steel_zone.add_continuous_addition(ContinuousGasAddition({"02": 100}, 1,
                                                                rate_unit=GasRateUnit.NORM_
→CUBIC_METER_PER_SEC))
    calc.with_reaction_zone(ReactionZone(area=10.0,
                                           left_zone=steel_zone, mass_transfer_coefficient_
\rightarrow left=1.0e-5.
                                           right_zone=slag_zone, mass_transfer_coefficient_
\rightarrowright=1.0e-6))
    result = calc.calculate()
    print(f"Stable phases in the steel melt: {result.get_stable_phases(steel_zone)}")
    print(f"C-content in steel vs. time: {result.get_composition_of_phase_group(steel_
\rightarrow zone,
                                                                                    ب
→ PhaseGroup.ALL_METAL) ['C'] }")
```

### 4.3.11 Additive Manufacturing calculations

For Additive Manufacturing calculations, everything that you can configure in the *AM Calculator* in Graphical Mode can also be configured in this calculation. In the minimal case, you do not need to specify any options (the material used by default is *IN625*).

```
from tc_python import *
with TCPython() as start:
    am_result = (start.with_additive_manufacturing()
                .with_steady_state_calculation()
                .calculate())
    print(f"Melt pool depth: {am_result.get_meltpool_depth()}")
    plotter, mesh = am_result.get_pyvista_plotter()
    plotter.add_mesh(mesh)
    plotter.show()
```

With several custom settings a basic calculation looks like this:

```
from tc_python import *
with TCPython() as start:
    am_result = (start.with_additive_manufacturing()
                  .with_steady_state_calculation()
                  .disable_fluid_flow_marangoni()
                  .disable_separate_materials()
                  .with_material_properties(MaterialProperties.from_library("IN718"))
                  .set_powder_density(90)
                  .set_ambient_temperature(15 + 273.15)
                  .set_base_plate_temperature(25 + 273.15)
                  .with_top_boundary_conditions(TopBoundaryConditions().set_radiation_
\rightarrow \text{emissivity}(0.2))
                  .with_heat_source(HeatSource.conical()
                                    .set_power(210)
                                     .set_absorptivity(80.0))
                  .calculate())
    print(f"Melt pool depth: {am_result.get_meltpool_depth()}")
    plotter, mesh = am_result.get_pyvista_plotter()
    plotter.add_mesh(mesh)
    plotter.show()
```

**Tip:** The graphical visualization and extraction of data from the 3D result dataset requires the package *pyvista*. More details are provided in the section *Optional packages*.

# 4.4 Result

All calculations have a method called calculate() that starts the calculations and when finished, returns a **Result**.

The Result classes have very different methods, depending on the type of calculation.

The **Result** is used to get numerical values from a calculation that has run.

The **Result** can be saved to disk by the method save\_to\_disk().

Previously saved results can be loaded by the method load\_result\_from\_disk() on the SetUp class.

#### **Example:**

```
# code above sets up the calculation
r = calculation.calculate()
time, meanRadius = r.get_mean_radius_of("AL3SC")
```

The **Result** objects are completely independent from calculations done before or after they are created. The objects return valid values corresponding to the calculation they were created from, for their lifetime. The only exception is if you call calculate() and not calculate\_with\_state() on a single equilibrium calculation.

As in the following example you can mix different calculations and results, and use old results after another calculation has run.

```
# ...
# some code to set up a single equilibrium calculation
# ...
single_eq_result = single_eq_calculation.calculate_with_state()
# ...
# some code to set up a precipitation calculation
# ...
prec_result = precipitation_calculation.calculate()
# ...
# some code to set up a Scheil calculation
# ...
scheil_result = scheil_calculations.calculate()
# now it is possible to get results from the single equilibrium calculation,
# without having to re-run it (because it has been calculated with saving of the state)
gibbs = single_eq_result.get_value_of("G")
```

### 4.4.1 DiffusionResult

The DiffusionResult class, that is returned when calling calculate() on any DiffusionCalculation, has the possibility to create a ContinuedDiffusionCalculation, in addition to the "normal" functionality for results. This makes it possible to run a diffusion calculation and then, depending on the result, change some settings and continue.

# 4.5 Property Model Framework

The *Python Property Model SDK* extends the Thermo-Calc software to enable you to create your own Property Models. A *Property Model* is a Python-based calculation that can use any TC-Python functionality (including diffusion and precipitation calculations) but is usable through the Graphical User Interface (UI) of Thermo-Calc in a more simple way. It is typically used to model material properties but by no means limited to that. Examples of Property Models provided by Thermo-Calc include Martensite and Pearlite formation in steel.

The Property Model Framework uses standard Python 3 beginning with Thermo-Calc 2021a and can access all TC-Python functionality and any Python package including *numpy*, *scipy*, *tensorflow*, etc. The actual calculation code is nearly identical, regardless if called from within a Property Model or from standard Python.

This is a complete rewrite of the original version of the framework that was based on Jython 2.7 and therefore had a number of limitations. **Property models written with the old Property Model Framework before Thermo-Calc 2021a are not compatible with the new framework**. However, the migration should be relatively easy because the syntax was changed as little as possible.

### 4.5.1 Property models vs. TC-Python

The main difference between a *Property Model* and regular *TC-Python code* is that a Property Model is directly integrated into the UI of Thermo-Calc via a plugin architecture while TC-Python code can only be accessed by programs and scripts written in Python.

The user should develop a Property Model if the functionality needs to be available from the Thermo-Calc UI, especially if it should be applied by other users not familiar to programming languages. Otherwise it is preferable to implement the functionality directly in a TC-Python program. If required, Property Models can as well be accessed from within TC-Python.

### 4.5.2 Architecture

Every Property Model needs to contain a class that implements the interface tc\_python.propertymodel\_sdk. PropertyModel. There are naming conventions that must to be fulfilled: the file name is required to follow the the pattern XYPythonModel.py and the name of the class needs to match this. Additionally the file must be placed in a directory named XYPython within the Property Model directory. The content of the placeholder XY can be freely chosen.

A simple complete Property Model, saved in a file called SimplePythonModel.py in the directory SimplePython, looks like this:

```
from tc_python import *

class SimplePythonModel(PropertyModel):
    def provide_model_category(self) -> List[str]:
        return ["Demo"]

    def provide_model_name(self) -> str:
        return "My Demo Model"

    def provide_model_description(self) -> str:
        return "This is a demo model."

    def provide_ui_panel_components(self) -> List[UIComponent]:
```

```
return [UIBooleanComponent("CHECKBOX", "Should this be checked?", "Simple_
-,checkbox", setting=False)]
def provide_calculation_result_quantities(self) -> List[ResultQuantity]:
    return [create_general_quantity("RESULT", "A result")]
def evaluate_model(self, context: CalculationContext):
    if context.get_ui_boolean_value("CHECKBOX"):
        self.logger.info("The checkbox is checked")
    # obtain the entered values from the GUI
    composition_as_mass_fraction = context.get_mass_fractions()
    temp_in_k = context.get_temperature()
    calc = context.system.with_single_equilibrium_calculation()
    # continue with a TC-Python calculation now ...
    context.set_result_quantity_value("RESULT", 5.0) # the value would normally_
-,have been calculated
```

The basic building blocks of the Property Model API are:

- *tc\_python.propertymodel\_sdk.ResultQuantity*: Defines a calculation result of a Property Model that will be provided to the UI after each model evaluation
- tc\_python.propertymodel\_sdk.CalculationContext: Provides access to the data from the UI (such as the entered composition and temperature) and to the current TC-Python system object which is the entrypoint for using TC-Python from within the Property Model
- tc\_python.propertymodel\_sdk.UIComponent: These are the UI-components that create the user interface of the Property Model within the model panel of the Thermo-Calc application UI. Different components are available (for example checkboxes, text fields and lists).

### 4.5.3 Property Model directory

The Property Model *py*-files need to be located within subdirectories of the *Property Model directory*, e.g. **PropertyModels/XYPython/XYPythonModel.py**. The default Property Model directory can be changed in the menu *Tools* -> *Options* in the graphical user interface.

Operating sys-	Default Property Model directory
tem	
Windows	C:\Program Files\Thermo-Calc\2025b\PropertyModels
Linux	/home/UserName/Thermo-Calc/2025b/PropertyModels or: /opt/Thermo-Calc/
	2025b/PropertyModels
MacOS	<pre>/Applications/Thermo-Calc-2025b.app/Contents/Resources/PropertyModels</pre>

Note: The Property Model directory location has been changed in the release 2023b.

# **BEST PRACTICES**

# 5.1 Re-use of the single equilibrium calculation state

The Thermo-Calc core keeps an internal state containing the data from previously performed calculations (such as composition of sublattices, previously formed phases, ...). This will be used for start values of future calculations (if not explicitly overwritten) and can strongly influence their convergence and calculation time. It can be useful to save and restore later the core-state **in advanced use cases**, these include:

- Improving the convergence speed in case of very complicated equilibria if a similar equilibrium had been calculated already before. Similarity refers here primarily to composition, temperature and entered phase set. This case can occur for example with the Nickel-database TCNi.
- Convenient and fast switching between states that have changed a lot (for example regarding suspended phases, numerical settings, ...)

The mechanism of saving and restoring the state is called bookmarking and is controlled with the two methods tc\_python.single\_equilibrium.SingleEquilibriumCalculation.bookmark\_state() and tc\_python.single\_equilibriumCalculation.set\_state\_to\_bookmark(). The following short example demonstrates how to switch between two different states easily in practice:

```
from tc_python import *
with TCPython() as session:
    calc = (session.
            select_database_and_elements("FEDEMO", ["Fe", "C"]).
            get_system().
            with_single_equilibrium_calculation().
            set_condition(ThermodynamicQuantity.temperature(), 2000.0).
            set_condition("X(C)", 0.01))
    calc.calculate()
   bookmark_temp_condition = calc.bookmark_state()
   calc.set_phase_to_fixed("BCC", 0.5)
    calc.remove_condition(ThermodynamicQuantity.temperature())
   bookmark_fixed_phase_condition = calc.bookmark_state()
   result_temp = calc.set_state_to_bookmark(bookmark_temp_condition)
   print("Conditions do contain temperature: {}".format(result_temp.get_conditions()))
    # this calculation had already been performed
   print("Stable phases (do not contain BCC): {}".format(result_temp.get_stable_
→phases()))
```

### 5.2 Re-use and saving of results

Before a calculation is run in TC-Python, a check is made to see if the exact same calculation has run before, and if that is the case, the result from the calculation can be loaded from disk instead of being re-calculated.

This functionality is always enabled within a script running TC-Python, but you can make it work the same way when re-running a script, or even when running a completely different script.

To use results from previous calculations, set a folder where TC-Python saves results, and looks for previous results.

This is controlled by the method tc\_python.server.SetUp.set\_cache\_folder().

```
from tc_python import *
with TCPython() as start:
    start.set_cache_folder("cache")
```

This folder can be a network folder and shared by many users. The calculation is not re-run if there is a previous TC-Python calculation with the same cache folder and exactly the same settings; the result is instead loaded from disk.

Another possibility is to explicitly save the result to disk and reload it later:

```
from tc_python import *
with TCPython() as start:
    # ... the system and calculator are set up and the calculation is performed
    result = calculator.calculate()
    result.save_to_disk("./result_dir")
```

You can then load the result again in another session:

```
from tc_python import *
with TCPython() as start:
    result = SetUp().load_result_from_disk().diffusion("./result_dir")
    x, frac = result.get_mole_fraction_of_component_at_time("Cr", 1000.0)
```

# 5.3 All TC-Python objects are non-copyable

*Never create a copy* of an instance of a class in TC-Python, neither by using the Python built-in function deepcopy() nor in any other way. All classes in TC-Python are proxies for classes in the underlying calculation server and normally hold references to result files. A copied class object in Python would consequently point to the same classes and result files in the calculation server.

Instead of making a copy, always create a new instance:

```
from tc_python import *
with TCPython() as start:
    system = start.select_database_and_elements("FEDEMO", ["Fe", "Cr"]).get_system()
    calculator = system.with_single_equilibrium_calculation()
    # *do not* copy the `calculator` object, create another one instead
    calculator_2 = system.with_single_equilibrium_calculation()
    # now you can use both calculators for different calculations ...
```

# 5.4 Python Virtual Environments

A Python installation can have several virtual environments. You can think of a virtual environment as a collection of third party packages that you have access to in your Python scripts. tc\_python is such a package.

To run TC-Python, you need to **install it into the same virtual environment** as your Python scripts are running in. If your scripts fail on import tc\_python, you need to execute the following command in the terminal of the same **Python environment** as your script is running in:

pip install TC\_Python-<version>-py3-none-any.whl

If you use the PyCharm IDE, you should do that within the **Terminal** built into the IDE. This **Terminal** runs automatically within your actual (virtual) environment.

To prevent confusion, it is recommend in most cases to *install TC-Python within your global interpreter*, for example by running the pip install command within your default Anaconda prompt.

# 5.5 Using with TCPython() efficiently

Normally you should call with TCPython() only once within each process.

**Note:** When leaving the *with*-clause, the Java backend engine process is stopped and all temporary data is deleted. Finally when entering the next *with*-clause a new Java process is started. This can take several seconds.

If appropriate, it is safe to run *with TCPython()* in a loop. Due to the time it takes this only makes sense if the calculation time per iteration is longer than a minute.

To prevent calling with TCPython() multiple times and cleaning up temporary data, you can use the following pattern.

```
from tc_python import *
# ...
def calculation(calculator):
    # you could also pass the `session` or `system` object if more appropriate
    calculator.set_condition("W(Cr)", 0.1)
    # further configuration ...
    result = calculator.calculate()
    # ...
    result.invalidate() # if the temporary data needs to be cleaned up immediately
if __name__ == '__main__':
    with TCPython() as session:
        system = session.select_database_and_elements("FEDEMO", ["Fe", "Cr"]).get_
\rightarrow system()
        calculator = system.with_single_equilibrium_calculation()
        for i in range(50):
            calculation(calculator)
```

# 5.6 Parallel calculations

It is possible to perform parallel calculations with TC-Python using multi-processing.

**Note:** Please note that **multi-threading is not suitable** for parallelization of computationally intensive tasks in Python. Additionally the Thermo-Calc core is not thread-safe. Using suitable Python-frameworks it is also possible to dispatch the calculations on different computers of a cluster.

A general pattern that can be applied is shown below. This code snippet shows how to perform single equilibrium calculations for different compositions in parallel. In the same way all other calculators of Thermo-Calc can be used or combined. For performance reasons in a real application, probably *numpy* arrays instead of Python arrays should be used.

Example:

```
import concurrent.futures
from tc_python import *

def do_perform(parameters):
    # this function runs within an own process
    with TCPython() as start:
        elements = ["Fe", "Cr", "Ni", "C"]
        calculation = (start.select_database_and_elements("FEDEMO", elements).
            get_system().
            with_single_equilibrium_calculation().
```
(continued from previous page)

```
set_condition("T", 1100).
                       set_condition("W(C)", 0.1 / 100).
                       set_condition("W(Ni)", 2.0 / 100))
        phase_fractions = []
        cr_contents = range(parameters["cr_min"],
                            parameters["cr_max"],
                            parameters["delta_cr"])
        for cr in cr_contents:
            result = (calculation.
                      set_condition("W(Cr)", cr / 100).
                      calculate())
            phase_fractions.append(result.get_value_of("NPM(BCC_A2)"))
   return phase_fractions
if __name__ == "__main__":
   parameters = [
        {"index": 0, "cr_min": 10, "cr_max": 15, "delta_cr": 1},
        {"index": 1, "cr_min": 15, "cr_max": 20, "delta_cr": 1}
   1
   bcc_phase_fraction = []
   num_processes = 2
   with concurrent.futures.ProcessPoolExecutor(num_processes) as executor:
        for result_from_process in zip(parameters, executor.map(do_perform, parameters)):
            # params can be used to identify the process and its parameters
            params, phase_fractions_from_process = result_from_process
            bcc_phase_fraction.extend(phase_fractions_from_process)
    # use the result in `bcc_phase_fraction`, for example for plotting
```

# 5.7 Handling crashes of the calculation engine

In some cases the Thermo-Calc calculation engine can crash. If batch calculations are performed, this brings down the complete batch. To handle this situation there is an exception you can use.

UnrecoverableCalculationException

That exception is thrown if the calculation server enters a state where no further calculations are possible. You should catch that exception outside of the *with TCPython()* clause and continue within a new *with*-clause.

Example:

```
from tc_python import *
for temperature in range(900, 1100, 10):
    try:
```

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```
with TCPython() as start:
            diffusion_result = (
                start.
                    select_thermodynamic_and_kinetic_databases_with_elements("FEDEMO",
\rightarrow "MFEDEMO", ["Fe", "Ni"]).
                    get_system().
                    with_isothermal_diffusion_calculation().
                        set_temperature(temperature).
                        set_simulation_time(108000.0).
                        add_region(Region("Austenite").
                            set_width(1E-4).
                            with_grid(CalculatedGrid.linear().set_no_of_points(50)).
                            with_composition_profile(CompositionProfile().
                                add("Ni", ElementProfile.linear(10.0, 50.0))
                            ).
                        add_phase("FCC_A1")).
                calculate())
            distance, ni_fraction = diffusion_result.get_mass_fraction_of_component_at_

→time("Ni", 108000.0)

            print(ni_fraction)
   except UnrecoverableCalculationException as e:
       print('Could not calculate. Continuing with next...')
```

# 5.8 Using TC-Python within a Jupyter Notebook or the Python console

TC-Python can also be used from within an interactive Jupyter Notebook and a Python console as well as similar products. The main difference from a regular Python program is that it is not recommended to use a *with*-clause to manage the TC-Python resources. That is only possible within a single Jupyter Notebook cell. Instead the standalone functions  $tc_python.server.start_api_server()$  and  $tc_python.server.stop_api_server()$  should be used for manually managing the resources.

**Note:** The *resources* of TC-Python are primarily the Java-process running on the backend side that performs the actual calculations and the temporary-directory of TC-Python that can grow to a large size over time, especially if precipitation calculations are performed. If a *with*-clause is used, these resources are automatically cleared after use.

You need to make sure that you execute the two functions tc\_python.server.start\_api\_server() and tc\_python.server.stop\_api\_server() exactly once within the Jupyter Notebook session. If not stopping TC-Python, extra Java-processes might be present and the temporary disk-space is not cleared. However, these issues can be resolved manually.

The temporary directories of TC-Python are named, for example, TC\_TMP4747588488953835507 that has a random ID. The temporary directory on different operating systems varies according to the pattern shown in the table.

Operating	sys-	Temporary directory
tem		
Windows		C:\Users{UserName}\AppData\Local\Temp\TC_TMP4747588488953835507
MacOS		/var/folders/g7/7du81ti_b7mm84n184fn3k910000lg/T/
		TC_TMP4747588488953835507
Linux		/tmp/TC_TMP4747588488953835507

In a Jupyter Notebook some features of an IDE such as auto-completion (*TAB-key*), available method lookup (press . and then *TAB*) and parameter lookup (set the cursor within the method-parenthesis and press *SHIFT* + *TAB* or *SHIFT* + *TAB* + *TAB* + *TAB* for the whole docstring) are also available.

# Example using TC-Python with a Jupyter Notebook:

💭 Jupyte	r TC-Python with Jupyter Notebook Last Checkpoint: a few seconds ago (autosaved)		Cogout
File Edit	View Insert Cell Kernel Widgets Help	rusted	Python [conda env:Anaconda3] O
+ %			
In [1]:	<pre>from tc_python import *</pre>		
In [2]:	<pre>start_api_server()</pre>		
In [3]:	<pre>system = SetUp().select_database_and_elements("FEDEMO", ["Fe", "Ni", "Cr"]).get_system() calc = system.with_single_equilibrium_calculation()</pre>		
In [4]:	temp = 825 # in K ni_conc = 10.0 # in wt-% cr_conc = 8.0 # in wt-%		
	<pre>calc. \     set_condition(ThermodynamicQuantity.temperature(), temp). \     set_condition(ThermodynamicQuantity.mass_fraction_of_a_component("Ni"), ni_conc / 100     set_condition(ThermodynamicQuantity.mass_fraction_of_a_component("Cr"), cr_conc / 100     result = calc.calculate()</pre>	0). \ 0)	
In [5]:	result.get_value_of(ThermodynamicQuantity.mole_fraction_of_a_phase("FCC_A1"))		
Out[5]:	0.3345580340424432		
In [6]:	<pre>stop_api_server()</pre>		

# 5.9 Property Model Framework

# 5.9.1 Debugging Property Model code

You can debug property models while running them from Thermo-Calc.

- Start Thermo-Calc and create a Property Model calculator.
- Select the model you want to debug and check the debug checkbox in the lower right corner of the Python code tab.

Dr	iving force					
Co	onfiguration	Description	Python code			
74				normatized_nriving_rorce)		
75	ging mode	ι:				
76	<pre>c_python.implementationpropertymodel_sdk import _PropertyModelProxy</pre>					
77	rtyModelProxy(model=DrivingForcePythonModel(), java_port=57334, python_port=65247)					
		Ŭ				
				C Debug		

Now the model that you want to debug has been updated with code needed to connect with Thermo-Calc.

• Start debugging the model in the IDE of your choice.

Note: You must use a Python interpreter where TC-Python is installed.

# In PyCharm it looks like this:

- [	ė									
pe	1 2		fro	m tc.	_python imp	ort *				
			¢cla	ss Di	rivingForce	PythonModel(F	PropertyModel):			
				RES	ULT_QUANTIT	<b>Y =</b> "drivingF				
	6 NORMALIZED_RESULT_QUANTITY = "normalizedDrivingForce"									
	7 MATRIX_PHASE_LIST = "matrix"									
				PRE	CIPITATE_PH	ASE_LIST = "p	precipitate"			
							Show Context Actions	Alt+Enter		
				def	provide_ca	lculation_re	🗂 <u>P</u> aste	Ctrl+V	ntity]:	
					results =	[create_ener	Copy / Paste Special		ULT_QUANTITY,	
							Column Selection <u>M</u> ode	Alt+Shift+Insert	uantity.text" <b>)),</b>	
						create_gene	Find <u>U</u> sages	Alt+F7	RMALIZED_RESULT_QUANTITY,	
							<u>R</u> efactor		izedResultQuantity.text"))]	
					return res	ULTS	Folding			
				dof	nnovide mo	del category				
				uer	return [se	lf get pror	Generate	Alt+Insert		
						erget_prot	Run 'DrivingForcePythonMo'	Ctrl+Shift+F10		
				def	provide_mo	del name(sel				
	~ -		I	1			🕻 Run 'DrivingForcePythonMo' wit	h Co <u>v</u> erage		

Note: When your IDE and Thermo-Calc have sucessfully connected, you will see this in the Thermo-Calc log:

You can stop the debug session in your IDE, change the model code, and start debugging again. The changes you made will take effect in Thermo-Calc without the need to restart. If you for instance changed the method evaluate\_model(), the change will take effect the next time you press *Perform*.

It is also possible to start the models from TC-Python. The workflow is exactly the same as described above, except instead of starting Thermo-Calc graphical user interface, you start a Python script and use the parameter debug\_model=True when selecting your model.

```
from tc_python import *
with TCPython() as start:
    property_model = (
        start.
            select_database_and_elements("FEDEMO", ["Fe", "C"]).
            get_system().
            with_property_model_calculation("my own Driving Force", debug_model=True).
            set_composition("C", 1.0).
            )
        property_model.calculate()
        ...
```

# 5.9.2 Developing Property Models in several files

You can split your Property Model code in several .py files, and there are two ways of doing that:

- side-by-side modules
- common modules

Side-by-side modules are Python files located in the same folder as the Property Model.

*Common* modules are Python files located in a folder outside of the Property Model folder, which makes it possible to share them with several models as a common library.

# 5.9.2.1 side-by-side modules

You are required to:

- Add a \_\_init\_\_.py file to your Property Model folder
- Add all imports of side-by-side modules in your main Property Model Python file also to the \_\_init\_\_.py file

#### Example:

CriticalTemperaturesPythonModel.py (The main Property Model file):

```
from CriticalTemperaturesPython import CriticalTemperatures
from tc_python import *
import numpy as np
class CriticalTemperaturesPythonModel(PropertyModel):
....
```

\_\_init\_\_.py:

from CriticalTemperaturesPython.critical\_temperatures\_library import CriticalTemperatures

If you are using PyCharm, the package name of the Property Model might be highlighted as an error, in this case you can mark the Property Model directory (i.e. the root of the present model directory) by right-clicking on it in the project window of PyCharm and marking it as *Sources Root*:



critical\_temperatures\_library.py:

```
from tc_python import *
import numpy as np
from scipy import optimize
from enum import Enum
class CriticalTemperatures(object):
....
```

**Note:** Modules installed in the Python interpreter such as *numpy*, *scipy*, etc can be imported as normal. This only concerns files imported as *side-by-side* modules.

# 5.9.2.2 common modules

*common* modules work very similar to *side-by-side* modules except the import statements are done in the "main" \_\_init\_\_.py file in *Property Model directory*.

You are required to:

- Add a \_\_\_init\_\_\_.py file to your property model folder.
- Add all imports of *common* modules in your main property model python file also to both the \_\_init\_\_.py file in *Property Model directory* AND the \_\_init\_\_.py of the property model.

# Example:

CriticalTemperaturesPythonModel.py (The main Property Model file):

from PropertyModels import Martensite
from tc\_python import \*

class CriticalTemperaturesPythonModel(PropertyModel):

...

\_\_init\_\_.py: (The init file located in the property model folder)

from PropertyModels import Martensite

\_\_init\_\_.py: (The init file located in *Property Model directory*)

from PropertyModels.common.martensite\_library import Martensite

The file critical\_temperatures\_library.py should in this example be located in a folder called common in the *Property Model directory*.

critical\_temperatures\_library.py:

```
from tc_python import *
import numpy as np
from scipy import optimize
from enum import Enum
class CriticalTemperatures(object):
....
```

Note: common modules must be located in folder called common.

# 5.9.3 Alternative Python for Property Models

# 5.9.3.1 Default bundled Python interpreter

Thermo-Calc is by default using a Python 3 interpreter bundled to the software for running the property models. It is containing the following major packages:

**Warning:** Any changes to the interpreter packages can therefore break Thermo-Calc and should be avoided. If the installation gets broken, it can be fixed by reinstalling Thermo-Calc after having removed it.

Please contact the Thermo-Calc support if you think that further packages might be useful in future releases. If these packages are insufficient for you, it is possible to use another Python-interpreter: *Configuring another Python interpreter*.

The interpreter is located in different places depending on the platform:

Operating sys-	Path to the bundled Python-interpreter
tem	
Windows	C:\Program Files\Thermo-Calc\2025b\python\python.exe
Linux	<pre>/home/UserName/Thermo-Calc/2025b/python/bin/python3</pre>
MacOS	/Applications/Thermo-Calc-2025b.app/Contents/Resources/python/
	bin/python3

# 5.9.3.2 Configuring another Python interpreter

If you require additional Python-packages or prefer to use your own interpreter installed on your system, you can change the interpreter used by Thermo-Calc to run the property models. Select **Tools** $\rightarrow$ **Options** in the Thermo-Calc GUI and modify the path to that of your Python 3 interpreter of choice:

Property model Python interpreter:         C:\Program Files\Thermo-Calc\2021a\python\python.exe         Mode	odify
--	-------

# 5.10 Process Metallurgy Calculations

# 5.10.1 Equilibrium calculations with changing elements between calculations

It is possible to add, change or remove additions after performing an equilibrium calculation using tc\_python. process\_metallurgy.equilibrium.EquilibriumCalculation.calculate(). This will change the elements being present in the system if the elements of the additions are differing. The Process Metallurgy Module will handle this situation by reloading the database with the latest set of elements. While this is an appropriate approach in most cases, there can be some disadvantages: reloading the database takes some time and the internal engine state is lost, which may lead to successive calculations failures in some situations.

To avoid the database reload, it is possible to add the respective elements to additions being present in all calculations (with a zero-fraction):

```
from tc_python import *
with TCPython() as session:
    calc = session.with_metallurgy().with_adiabatic_equilibrium_
→calculation(ProcessDatabase.OXDEMO)
    # add the element Al with zero-fraction already
    steel = EquilibriumAddition({'Fe': None, 'C': 4, 'Al': 0}, amount=100.0e3,...
\rightarrow temperature=1700 + 273.15)
    slag = EquilibriumAddition({'Ca0': 70, 'Si02': 30}, amount=3.0e3, temperature=1700 +_
\leftrightarrow 273.15)
    al_addition = EquilibriumAddition({'Al': 100}, amount=1.0e3)
    (calc
     .add addition(steel)
     .add_addition(slag))
    result_1 = calc.calculate()
    calc.add_addition(al_addition)
    result_2 = calc.calculate()
    # evaluate the result as required ...
```

Or to add a later addition already before the first call to calculate() with a zero amount:

from tc\_python import \*
with TCPython() as session:

(continues on next page)

(continued from previous page)

```
calc = session.with_metallurgy().with_adiabatic_equilibrium_

→calculation(ProcessDatabase.OXDEMO)

   steel = EquilibriumAddition({'Fe': None, 'C': 4}, amount=100.0e3, temperature=1700 +_
\leftrightarrow 273.15)
   slag = EquilibriumAddition({'Ca0': 70, 'Si02': 30}, amount=3.0e3, temperature=1700 +_
\rightarrow 273.15)
   # add the addition for now with zero-amount
   al_addition = EquilibriumAddition({'Al': 100}, amount=0)
   (calc
     .add_addition(al_addition)
     .add_addition(steel)
    .add_addition(slag))
   result_1 = calc.calculate()
   calc.update_addition(al_addition.set_amount(1.0e3))
   result_2 = calc.calculate()
   # evaluate the result as required ...
```

# 5.10.2 Zones

TC-Python is providing a *framework* for building time-dependent kinetic simulations of industrial and academic metallurgical processes where liquid phases are important. It is based on an *Effective Equilibrium Reaction Zone* (EERZ) approach which is separating a process into different zones. These zones have identical temperature and composition and are called *bulk zones*. Such zones can be in contact and react with each other by *reaction zones*. That means a *reaction zone* is modelling the interface between two *bulk zones*. One bulk zone is typically the steel melt and another bulk zone the top slag.

# 5.10.3 Applications

While this approach can in principle be extended to any number of zones, in the current release TC-Python is providing **only one reaction zone**. Practical work has however proven that this limitation is not critical for a lot of industrial processes, including ladle furnaces, AOD- and VOD-converters. Even more processes can be modelled with some limit of accuracy.

The reason for the power of the current implementation is that a number of important process features can be included:

- heating (tc\_python.process\_metallurgy.process.Zone.add\_power())
- cooling (tc\_python.process\_metallurgy.process.Zone.add\_power())
- heat transfer between bulk zones (tc\_python.process\_metallurgy.process.ReactionZone. add\_heat\_transfer())
- · inclusion formation
- inclusion flotation and other transfer of phase groups between bulk zones (tc\_python.process\_metallurgy. process.ReactionZone.add\_transfer\_of\_phase\_group())

- addition of material and gas at any time in any zone (tc\_python.process\_metallurgy.process.Zone. add\_addition() / tc\_python.process\_metallurgy.process.Zone.add\_continuous\_addition())
- an exhaust gas zone collecting all formed gas (tc\_python.process\_metallurgy.process. ProcessSimulationResult.get\_exhaust\_gas())
- time-dependent definition of most parameters (e.g., mass transfer coefficient, transfer of phase group, heating, etc.)

Please note that many of these features are called as well a *reaction zone* in other EERZ model implementations.

# 5.10.4 Implementation of practical process models

The Process Metallurgy Module has been successfully applied to a number of industrial processes.

Due to the broad range of industrial metallurgical processes, TC-Python is not providing ready-to-use models for certain processes. There are however examples available for common processes and this collection will be extended over time. The implementation of a model is an abstraction of the real process and should always be kept as simple as possible. Practical experience has proven that in many situations not more than one *reaction zone* is required.

The mass transfer coefficient is a fundamental parameter describing the kinetics in a *reaction zone* and is generally an empirical parameter. It depends however mostly on the geometry and stirring conditions in the process and not on the material compositions. Further on, the mass transfer coefficient has usually typical values for a given process - regardless of the actual furnace. That means that existing suggestions from the literature can be used as a starting point to derive the actual mass transfer coefficient for the process of interest.

# CHAPTER

# SIX

# **API REFERENCE**

# 6.1 Calculations

# 6.1.1 Module "single\_equilibrium"

# class tc\_python.single\_equilibrium.AbstractSingleEquilibriumCalculation(calculator)

Bases: AbstractCalculation

Abstract configuration required for a single equilibrium calculation.

Note: This is an abstract class that cannot be used directly.

# disable\_global\_minimization()

Turns the global minimization completely off.

# Returns

This SingleEquilibriumCalculation object

# enable\_global\_minimization()

Turns the global minimization on (using the default settings).

# Returns

This SingleEquilibriumCalculation object

# $\texttt{get\_components()} \rightarrow List[str]$

Returns a list of components in the system (including all components auto-selected by the database(s)).

# Returns

The components

# $\texttt{get\_gibbs\_energy\_addition\_for}(\textit{phase: str}) \rightarrow \texttt{float}$

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

# **Parameters**

phase – Specify the name of the (stoichiometric or solution) phase with the addition

# Returns

Gibbs energy addition to G per mole formula unit.

#### $get_system_data() \rightarrow SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using *with\_system\_modifications()*.

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as \*.tdb-file.

#### Returns

The system data

#### run\_poly\_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

#### Parameters

command - The Thermo-Calc Console Mode command

# Returns

This SingleEquilibriumCalculation object

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

# set\_component\_to\_entered(component: str)

Sets the specified component to the status ENTERED, that is the default state.

# Parameters

component – The component name or ALL\_COMPONENTS

#### Returns

This SingleEquilibriumCalculation object

# set\_component\_to\_suspended(component: str, reset\_conditions: bool = False)

Sets the specified component to the status SUSPENDED, i.e. it is ignored in the calculation.

#### **Parameters**

- **reset\_conditions** if 'True' also remove composition conditions for the component if they are defined
- **component** The component name or *ALL\_COMPONENTS*

# Returns

This SingleEquilibriumCalculation object

# set\_gibbs\_energy\_addition\_for(phase: str, gibbs\_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs\_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

# Parameters

- phase Specify the name of the (stoichiometric or solution) phase with the addition
- gibbs\_energy Addition to G per mole formula unit

# Returns

This SingleEquilibriumCalculation object

#### set\_phase\_to\_dormant(phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

**Parameters** 

phase – The phase name or ALL\_PHASES for all phases

#### Returns

This SingleEquilibriumCalculation object

# set\_phase\_to\_entered(phase: str, amount: float = 1.0)

Sets the phase to the status ENTERED, that is the default state.

#### **Parameters**

- phase The phase name or ALL\_PHASES for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

#### Returns

This SingleEquilibriumCalculation object

# set\_phase\_to\_fixed(phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

# Parameters

- **phase** The phase name
- **amount** The fixed phase fraction (between 0.0 and 1.0)

#### Returns

This SingleEquilibriumCalculation object

# set\_phase\_to\_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

# phase – The phase name or ALL\_PHASES for all phases Returns

**Parameters** 

This SingleEquilibriumCalculation object

#### with\_options(options: SingleEquilibriumOptions)

Sets the simulation options.

#### Parameters

options – The simulation options

#### Returns

This SingleEquilibriumCalculation object

with\_reference\_state(component: str, phase: str = 'SER', temperature: float = -1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

#### Parameters

- component The name of the element must be given.
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT\_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The Pressure (in Pa) for the reference state.

#### Returns

#### This SingleEquilibriumCalculation object

# with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

**Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

#### **Parameters**

system\_modifications - The system modification to be performed

#### Returns

This SingleEquilibriumCalculation object

# class tc\_python.single\_equilibrium.SingleEquilibriumCalculation(calculator)

Bases: AbstractSingleEquilibriumCalculation

Configuration for a single equilibrium calculation.

Note: Specify the conditions and possibly other settings, the calculation is performed with calculate().

# **bookmark\_state**(*bookmark\_id:* $str = ") \rightarrow str$

Puts a "bookmark" on the current calculation-state of the calculator allowing the program to return to this state later as needed.

By bookmarking a state, you can simplify the convergence of equilibria when they strongly depend on the starting conditions (i.e. the state). Also use it to improve performance by running a calculation, then bookmarking it, and later returning to it for other equilibria whose conditions are "close" to the bookmarked equilibrium.

This method is used in combination with the method set\_state\_to\_bookmark().

#### Parameters

bookmark\_id – The bookmark id. If omitted a generated id is used and returned

# Returns

The bookmark id

# $calculate() \rightarrow SingleEquilibriumTempResult$

Performs the calculation and provides a temporary result object that is only valid until something gets changed in the calculation state. The method *calculate()* is the default approach and should be used in most cases.

# Returns

A new *SingleEquilibriumTempResult* object which can be used to get specific values from the calculated result. It is undefined behavior to use that object after the state of the calculation has been changed.

**Warning:** If the result object should be valid for the whole program lifetime, use calculate\_with\_state() instead.

# $calculate_with_state(timeout_in_minutes: float = 0.0) \rightarrow SingleEquilibriumResult$

Performs the calculation and provides a result object that reflects the present state of the calculation during the whole lifetime of the object.

**Note:** Because this method has performance and temporary disk space overhead (i.e. it is resource heavy), only use it when it is necessary to access the result object after the state is changed. In most cases you should use the method *calculate()*.

# Parameters

**timeout\_in\_minutes** – Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a Unre-coverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

#### Returns

A new *SingleEquilibriumResult* object which can be used later at any time to get specific values from the calculated result.

# disable\_global\_minimization()

Turns the global minimization completely off.

#### Returns

#### enable\_global\_minimization()

Turns the global minimization on (using the default settings).

# Returns

This SingleEquilibriumCalculation object

# $\texttt{get\_components()} \rightarrow List[str]$

Returns a list of components in the system (including all components auto-selected by the database(s)).

#### Returns

The components

# 

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

# Parameters

phase – Specify the name of the (stoichiometric or solution) phase with the addition

## Returns

Gibbs energy addition to G per mole formula unit.

```
get_interfacial_energy(matrix_phase: str, precipitate_phases: List[str], zero_volume_elements:
List[str] = ['C', 'N']) \rightarrow Dict[str, float]
```

Estimates the interfacial energy between a matrix phase and a precipitate phase using thermodynamic data from a CALPHAD database. The approximation model is based on Becker's bond energy approach.

Default: elements with no contribution to volume are C and N.

#### **Parameters**

- **matrix\_phase** The matrix phase.
- **precipitate\_phases** The list of precipitate phases for which interfacial energy between them and the matrix phase is to be calculated.
- zero\_volume\_elements The elements that are assumed to not contribute to the volume.

#### Returns

A dictionary containing interfacial energy per precipitate phase.

# $get_system_data() \rightarrow SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using *with\_system\_modifications()*.

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as \*.tdb-file.

# Returns

The system data

#### remove\_all\_conditions()

Removes all set conditions.

#### Returns

#### remove\_condition(quantity: Union[ThermodynamicQuantity, str])

Removes the specified condition.

#### **Parameters**

**quantity** – the thermodynamic quantity to set as condition; a Console Mode syntax string can be used as an alternative (for example "X(Cr)")

#### Returns

This SingleEquilibriumCalculation object

#### run\_poly\_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

# Parameters

command – The Thermo-Calc Console Mode command

#### Returns

This SingleEquilibriumCalculation object

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

# set\_component\_to\_entered(component: str)

Sets the specified component to the status ENTERED, that is the default state.

#### Parameters

**component** – The component name or *ALL\_COMPONENTS* 

#### Returns

This SingleEquilibriumCalculation object

# set\_component\_to\_suspended(component: str, reset\_conditions: bool = False)

Sets the specified component to the status SUSPENDED, i.e. it is ignored in the calculation.

#### Parameters

- **reset\_conditions** if 'True' also remove composition conditions for the component if they are defined
- component The component name or ALL\_COMPONENTS

# Returns

This SingleEquilibriumCalculation object

# set\_condition(quantity: Union[ThermodynamicQuantity, str], value: float)

Sets the specified condition.

# **Parameters**

- **quantity** The thermodynamic quantity to set as condition; a Console Mode syntax string can be used as an alternative (for example "X(Cr)")
- value The value of the condition

#### Returns

# set\_gibbs\_energy\_addition\_for(phase: str, gibbs\_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs\_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

# Parameters

- phase Specify the name of the (stoichiometric or solution) phase with the addition
- gibbs\_energy Addition to G per mole formula unit

# Returns

This SingleEquilibriumCalculation object

# set\_phase\_to\_dormant(phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

Parameters phase – The phase name or *ALL\_PHASES* for all phases

# Returns

This SingleEquilibriumCalculation object

# set\_phase\_to\_entered(phase: str, amount: float = 1.0)

Sets the phase to the status ENTERED, that is the default state.

## **Parameters**

- phase The phase name or ALL\_PHASES for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

# Returns

This SingleEquilibriumCalculation object

# set\_phase\_to\_fixed(phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

# Parameters

• phase – The phase name

• **amount** – The fixed phase fraction (between 0.0 and 1.0)

#### Returns

This SingleEquilibriumCalculation object

# set\_phase\_to\_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

# Parameters

phase – The phase name or ALL\_PHASES for all phases

# Returns

#### **set\_state\_to\_bookmark**(bookmark\_id: str) → SingleEquilibriumTempResult

Resets the calculation state to a previously bookmarked state.

After calling this method, the calculation behaves exactly as it would after the bookmarked calculation ran.

This method is used in combination with the method *bookmark\_state()*.

# Parameters

**bookmark\_id** – The bookmark id of the state to return to.

# Returns

A new *SingleEquilibriumTempResult* object which can be used to get specific values from the calculated result. It is undefined behavior to use that object after the state of the calculation has been changed.

# with\_options(options: SingleEquilibriumOptions)

Sets the simulation options.

```
Parameters
options – The simulation options
```

Returns

This SingleEquilibriumCalculation object

# with\_reference\_state(component: str, phase: str = 'SER', temperature: float = -1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

#### Parameters

- component The name of the element must be given.
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT\_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The Pressure (in Pa) for the reference state.

#### Returns

## with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

**Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

#### Parameters

system\_modifications – The system modification to be performed

Returns

This SingleEquilibriumCalculation object

# class tc\_python.single\_equilibrium.SingleEquilibriumOptions

Bases: object

General simulation conditions for the thermodynamic calculations.

# disable\_approximate\_driving\_force\_for\_metastable\_phases()

Disables the approximation of the driving force for metastable phases.

Default: Enabled

**Note:** When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it is important that these driving forces are correct, use *disable\_approximate\_driving\_force\_for\_metastable\_pha* to force the calculation to converge for the metastable phases.

#### Returns

This SingleEquilibriumOptions object

#### disable\_control\_step\_size\_during\_minimization()

Disables stepsize control during minimization (non-global).

Default: Enabled

#### Returns

This SingleEquilibriumOptions object

# disable\_force\_positive\_definite\_phase\_hessian()

Disables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

Default: Enabled

# Returns

This SingleEquilibriumOptions object

## enable\_approximate\_driving\_force\_for\_metastable\_phases()

Enables the approximation of the driving force for metastable phases.

Default: Enabled

**Note:** When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it is important that these driving forces are correct, use *disable\_approximate\_driving\_force\_for\_metastable\_pha* to force the calculation to converge for the metastable phases.

#### Returns

This SingleEquilibriumOptions object

# enable\_control\_step\_size\_during\_minimization()

Enables stepsize control during normal minimization (non-global).

Default: Enabled

#### Returns

This SingleEquilibriumOptions object

# enable\_force\_positive\_definite\_phase\_hessian()

Enables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

Default: Enabled

#### Returns

This SingleEquilibriumOptions object

# set\_global\_minimization\_max\_grid\_points(max\_grid\_points: int = 2000)

Sets the maximum number of grid points in global minimization. Only applicable if global minimization is actually used.

Default: 2000 points

Parameters
max\_grid\_points – The maximum number of grid points

Returns

This SingleEquilibriumOptions object

### set\_max\_no\_of\_iterations(max\_no\_of\_iterations: int = 500)

Set the maximum number of iterations.

Default: max. 500 iterations

**Note:** As some models give computation times of more than 1 CPU second/iteration, this number is also used to check the CPU time and the calculation stops if 500 CPU seconds/iterations are used.

#### **Parameters**

max\_no\_of\_iterations - The max. number of iterations

#### Returns

This SingleEquilibriumOptions object

# set\_required\_accuracy(accuracy: float = 1e-06)

Sets the required relative accuracy.

Default: 1.0E-6

**Note:** This is a relative accuracy, and the program requires that the relative difference in each variable must be lower than this value before it has converged. A larger value normally means fewer iterations but less accurate solutions. The value should be at least one order of magnitude larger than the machine precision.

Parameters accuracy – The required relative accuracy

# Returns

This SingleEquilibriumOptions object

## set\_smallest\_fraction(smallest\_fraction: float = 1e-12)

Sets the smallest fraction for constituents that are unstable.

It is normally only in the gas phase that you can find such low fractions.

The **default value** for the smallest site-fractions is 1E-12 for all phases except for IDEAL phase with one sublattice site (such as the GAS mixture phase in many databases) for which the default value is always as 1E-30.

## **Parameters**

smallest\_fraction – The smallest fraction for constituents that are unstable

#### Returns

This SingleEquilibriumOptions object

#### class tc\_python.single\_equilibrium.SingleEquilibriumResult(result)

# Bases: AbstractResult

Result of a single equilibrium calculation, it can be evaluated using a Quantity or Console Mode syntax.

# change\_pressure(pressure: float)

Change the pressure and re-evaluate the results from the equilibrium without minimizing Gibbs energy, i.e. with higher performance. The properties are calculated at the new pressure using the phase amount, temperature and composition of phases from the initial equilibrium. Use *get\_value\_of()* to obtain them.

#### Parameters

pressure – The pressure [Pa]

Returns

This SingleEquilibriumCalculation object

#### change\_temperature(temperature: float)

Change the temperature and re-evaluate the results from the equilibrium without minimizing Gibbs energy, i.e. with high performance. The properties are calculated at the new temperature using the phase amount, pressure and composition of phases from the initial equilibrium. Use *get\_value\_of()* to obtain them.

**Note:** This is typically used when calculating room temperature properties (e.g. density) for a material when it is assumed that the equilibrium phase amount and composition freeze-in at a higher temperature during cooling.

#### Parameters

temperature – The temperature [K]

# Returns

This SingleEquilibriumCalculation object

# $\texttt{get\_components()} \rightarrow List[str]$

Returns the names of the components selected in the system (including any components auto-selected by the database(s)).

# Returns

The names of the selected components

# $\texttt{get\_conditions()} \rightarrow List[str]$

Returns the conditions.

# Returns

The selected conditions

#### $get_phases() \rightarrow List[str]$

Returns the phases present in the system due to its configuration. It also contains all phases that have been automatically added during the calculation, this is the difference to the method System. get\_phases\_in\_system().

## Returns

The names of the phases in the system including automatically added phases

# get\_stable\_phases() $\rightarrow$ List[str]

Returns the stable phases (i.e. the phases present in the current equilibrium).

## Returns

The names of the stable phases

# get\_value\_of(quantity: Union[ThermodynamicQuantity, str]) $\rightarrow$ float

Returns a value from a single equilibrium calculation.

# Parameters

**quantity** – The thermodynamic quantity to get the value of; a Console Mode syntax strings can be used as an alternative (for example "NPM(FCC\_A1)")

# Returns

The requested value

# run\_poly\_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine. This affects only the state of the result object.

#### Parameters

command - The Thermo-Calc Console Mode command

Returns

This SingleEquilibriumCalculation object

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

# save\_to\_disk(path: str)

Saves the result to disk. Note that the result is a folder, containing potentially many files. The result can later be loaded with load\_result\_from\_disk()

#### **Parameters**

**path** – the path to the folder you want the result to be saved in. It can be relative or absolute.

# Returns

this SingleEquilibriumResult object

# class tc\_python.single\_equilibrium.SingleEquilibriumTempResult(result)

# Bases: AbstractResult

Result of a single equilibrium calculation that is only valid until something gets changed in the calculation state. It can be evaluated using a Quantity or Console Mode syntax.

**Warning:** Note that it is undefined behavior to use that object after something has been changed in the state of the calculation, this will result in an InvalidResultStateException exception being raised.

#### change\_pressure(pressure: float)

Change the pressure and re-evaluate the results from the equilibrium without minimizing Gibbs energy, i.e. with higher performance. The properties are calculated at the new pressure using the phase amount, temperature and composition of phases from the initial equilibrium. Use *get\_value\_of()* to obtain them.

Parameters pressure – The pressure [Pa]

# Returns

This SingleEquilibriumCalculation object

# change\_temperature(temperature: float)

Change the temperature and re-evaluate the results from the equilibrium without minimizing Gibbs energy, i.e. with high performance. The properties are calculated at the new temperature using the phase amount, pressure and composition of phases from the initial equilibrium. Use *get\_value\_of()* to obtain them.

**Note:** This is typically used when calculating room temperature properties (e.g. density) for a material when it is assumed that the equilibrium phase amount and composition freeze-in at a higher temperature during cooling.

#### Parameters

temperature – The temperature [K]

#### Returns

This SingleEquilibriumCalculation object

# $\texttt{get\_components()} \rightarrow List[str]$

Returns the names of the components selected in the system (including any components auto-selected by the database(s)).

#### Returns

The names of the selected components

#### Raises

**InvalidResultStateException** – If something has been changed in the state of the calculation since that result object has been created

# $get\_conditions() \rightarrow List[str]$

Returns the conditions.

# Returns

List containing the selected conditions

#### Raises

**InvalidResultStateException** – If something has been changed in the state of the calculation since that result object has been created

# get\_phases() $\rightarrow$ List[str]

Returns the phases present in the system due to its configuration. It also contains all phases that have been automatically added during the calculation, this is the difference to the method System. get\_phases\_in\_system().

# Returns

The names of the phases in the system including automatically added phases

#### Raises

**InvalidResultStateException** – If something has been changed in the state of the calculation since that result object has been created

# get\_stable\_phases() $\rightarrow$ List[str]

Returns the stable phases (i.e. the phases present in the current equilibrium).

#### Returns

The names of the stable phases

#### Raises

*InvalidResultStateException* – If something has been changed in the state of the calculation since that result object has been created

get\_value\_of(quantity: Union[ThermodynamicQuantity, str])  $\rightarrow$  float

Returns a value from a single equilibrium calculation.

#### **Parameters**

**quantity** – The thermodynamic quantity to get the value of; a Console Mode syntax strings can be used as an alternative (for example "NPM(FCC\_A1)")

# Returns

The requested value

# Raises

**InvalidResultStateException** – If something has been changed in the state of the calculation since that result object has been created

# run\_poly\_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

#### Parameters

command - The Thermo-Calc Console Mode command

#### Returns

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

# 6.1.2 Module "batch\_equilibrium"

#### class tc\_python.batch\_equilibrium.BatchEquilibriumCalculation(calculator)

# Bases: AbstractCalculation

Configuration for a series of single equilibrium calculations performed in a vectorized fashion.

Note: Specify the conditions and call calculate().

**Tip:** The performance of batch equilibrium calculations can be significantly better than looping and using SingleEquilibriumCalculation if the actual Thermo-Calc calculation is fast. There is little advantage if the Thermo-Calc equilibrium calculations take a long time (typically for large systems and databases).

# **calculate**(*quantities: List[Union[ThermodynamicQuantity, str]], logging\_frequency: int = 10, timeout in minutes: float = 0.0*) $\rightarrow$ *BatchEquilibriumResult*

Runs the batch equilibrium calculation. The calculated *BatchEquilibriumResult* can then be queried for the values of the quantities specified.

# Example:

>>> quantities = ['G', 'X(BCC)']

#### **Parameters**

- quantities A list of the quantities to be calculated.
- **logging\_frequency** Determines how often logging should be done.
- **timeout\_in\_minutes** Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

#### Returns

A *BatchEquilibriumResult* which later can be used to get specific values from the calculated result.

# disable\_global\_minimization()

Turns the global minimization completely off.

#### Returns

This BatchEquilibriumCalculation object

# enable\_global\_minimization()

Turns the global minimization on (using the default settings).

#### Returns

This BatchEquilibriumCalculation object

# $get\_components() \rightarrow List[str]$

Returns a list of components in the system (including all components auto-selected by the database(s)).

#### Returns

The components

# 

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

# Parameters

phase – Specify the name of the (stoichiometric or solution) phase with the addition

#### Returns

Gibbs energy addition to G per mole formula unit.

# get\_system\_data() $\rightarrow$ SystemData

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using *with\_system\_modifications()*.

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as \*.tdb-file.

# Returns

The system data

# remove\_all\_conditions()

Removes all set conditions.

## Returns

This BatchEquilibriumCalculation object

#### remove\_condition(quantity: Union[ThermodynamicQuantity, str])

Removes the specified condition.

## Parameters

**quantity** – the thermodynamic quantity to set as condition; a Console Mode syntax string can be used as an alternative (for example "X(Cr)")

#### Returns

This BatchEquilibriumCalculation object

# run\_poly\_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

#### Parameters

command - The Thermo-Calc Console Mode command

#### Returns

This BatchEquilibriumCalculation object

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

#### set\_component\_to\_entered(component: str)

Sets the specified component to the status ENTERED, that is the default state.

#### **Parameters**

**component** – The component name or ALL\_COMPONENTS

#### Returns

This BatchEquilibriumCalculation object

## set\_component\_to\_suspended(component: str, reset\_conditions: bool = False)

Sets the specified component to the status SUSPENDED, i.e. it is ignored in the calculation.

#### Parameters

- reset\_conditions if 'True' also remove composition conditions for the component if they are defined
- **component** The component name or *ALL\_COMPONENTS*

#### Returns

This BatchEquilibriumCalculation object

set\_condition(quantity: Union[ThermodynamicQuantity, str], value: float)

Sets the specified condition.

# Parameters

- **quantity** The thermodynamic quantity to set as condition; a Console Mode syntax string can be used as an alternative (for example "X(Cr)")
- **value** The value of the condition

# Returns

This BatchEquilibriumCalculation object

Set the conditions of the equilibria to be calculated.

This is done by sending a list of equilibria at once.

Each equilibrium itself is a list of conditions that will be changed for that equilibrium.

A condition is described by a tuple containing:

- 1. A Console Mode syntax string or a ThermodynamicQuantity instance,
- 2. A float value specifying the value of the condition.

#### Example:

>>> [[('T', 800), ('X(Cr)', 0.1)], [('T', 850), ('X(Cr)', 0.11)]]

You can use ThermodynamicQuantity instead of a Console Mode syntax string when specifying type of condition.

Example:

#### Parameters

equilibria – The list of equilibria

Returns

This BatchEquilibriumCalculation object

#### set\_gibbs\_energy\_addition\_for(phase: str, gibbs\_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs\_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

#### **Parameters**

• phase - Specify the name of the (stoichiometric or solution) phase with the addition

• gibbs\_energy - Addition to G per mole formula unit

# Returns

This BatchEquilibriumCalculation object

#### set\_phase\_to\_dormant(phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

Parameters

phase – The phase name or ALL\_PHASES for all phases

#### Returns

This BatchEquilibriumCalculation object

## set\_phase\_to\_entered(phase: str, amount: float = 1.0)

Sets the phase to the status ENTERED, that is the default state.

#### **Parameters**

- phase The phase name or ALL\_PHASES for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

# Returns

This BatchEquilibriumCalculation object

# set\_phase\_to\_fixed(phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

#### Parameters

- **phase** The phase name
- **amount** The fixed phase fraction (between 0.0 and 1.0)

#### Returns

This BatchEquilibriumCalculation object

# set\_phase\_to\_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

#### Parameters

phase – The phase name or ALL\_PHASES for all phases

#### Returns

This BatchEquilibriumCalculation object

#### with\_options(options: SingleEquilibriumOptions)

Sets the simulation options.

# Parameters options – The simulation options

Returns

#### This BatchEquilibriumCalculation object

# with\_reference\_state(component: str, phase: str = 'SER', temperature: float = -1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

#### Parameters

- **component** The name of the element must be given.
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT\_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The Pressure (in Pa) for the reference state.

#### Returns

This BatchEquilibriumCalculation object

# with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

**Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

Parameters system\_modifications – The system modification to be performed

Returns

This BatchEquilibriumCalculation object

class tc\_python.batch\_equilibrium.BatchEquilibriumResult(result)

Bases: object

Result of a batch equilibrium calculation. This can be used to query for specific values.

**get\_values\_of**(*quantity: Union*[ThermodynamicQuantity, *str*]) → List[float]

Returns values from a batch equilibrium calculation.

**Warning:** The quantity must be one of the quantities specified for the *BatchEquilibriumCalculation* object that created the result object.

Example:

```
>>> batch_result = batch_calculation.calculate(quantities = ['G', 'X(BCC)'])
>>> batch_result.get_values_of('G')
```

#### Parameters

**quantity** – the thermodynamic quantity to get the value of; a Console Mode syntax strings can be used as an alternative (for example "NPM(FCC\_A1)")

# invalidate()

Invalidates the object and frees the disk space used by it.

**Note:** This is only required if the disk space occupied by the object needs to be released during the calculation. No data can be retrieved from the object afterwards.

# 6.1.3 Module "precipitation"

```
class tc_python.precipitation.FixedGrainSize(grain_radius: float = 0.0001)
```

Bases: GrainGrowthModel

```
set_grain_aspect_ratio(grain_aspect_ratio: float = 1.0)
```

Enter a numerical value. Default: 1.0.

**Parameters** grain\_aspect\_ratio – The grain aspect ratio [-]

## class tc\_python.precipitation.GrainGrowth(grain\_size\_distribution: GrainSizeDistribution)

Bases: GrainGrowthModel

#### disable\_zener\_pinning()

Disable Zener pinning to ignore the particle pinning effect on the grain growth. Zener pinning is by default disabled when no grain size distribution is defined, i.e. a single constant grain size is used. The setting is by default enabled when a grain size distribution is defined.

#### Returns

This GrainSizeDistribution object

# enable\_zener\_pinning()

Enable Zener pinning to simulate the particle pinning effect on the grain growth. The setting is by default enabled when a grain size distribution is defined.

# Returns

This GrainSizeDistribution object

#### set\_grain\_boundary\_energy(energy: float = 0.5)

Set the energy of the grain boundary.

#### Parameters

energy – The grain boundary energy [J/m2]

#### Returns

This GrainSizeDistribution object

# set\_grain\_boundary\_mobility\_activation\_energy(activation\_energy: float = 242000.0)

Set the grain boundary mobility activation energy where the mobility is defined by an Arrhenius type of equation.

#### Parameters

activation\_energy – The mobility activation energy [J/mol]

# Returns

This GrainSizeDistribution object

# set\_grain\_boundary\_mobility\_pre\_factor(pre\_factor: float = 0.004)

Set the grain boundary mobility prefactor where the mobility is defined by an Arrhenius type of equation.

#### Parameters

 $\ensuremath{\text{pre_factor}}\xspace$  – The grain boundary mobility pre factor  $[m^4/(J\ s)]$ 

# Returns

This GrainSizeDistribution object

## class tc\_python.precipitation.GrainGrowthModel

Bases: object

Factory class providing objects representing a grain growth model.

# classmethod fixed\_grain\_size(grain\_radius: float = 0.0001)

Fixed grain radius size. Default: 1.0E-4 m

#### Parameters

grain\_radius – The grain radius / size [m]

# classmethod grain\_growth(grain\_size\_distribution: GrainSizeDistribution)

Sets the initial grain size distribution for the matrix. **Default**: If the initial grain size distribution is not explicitly provided, a constant average grains size will be used and no grain growth evaluated during the simulation.

Tip: Use this option if you want to study the further evolution of an existing microstructure.

## Parameters

grain\_size\_distribution – grain size distribution

# class tc\_python.precipitation.GrainSizeDistribution

Bases: object

Represents the grain size distribution at a certain time.

add\_radius\_and\_number\_density(radius: float, number\_density: float)

Adds a radius and number density pair to the grain size distribution.

# Parameters

• radius – The radius [m]

• number\_density – The number of grains per unit volume per unit length [m^-4]

## Returns

This GrainSizeDistribution object

class tc\_python.precipitation.GrowthRateModel(value)

Bases: Enum

Choice of the used growth rate model for a precipitate.

The most efficient model is the *Simplified model*, which is the default and applicable to most alloy systems under the assumption that either the supersaturation is small, or the alloying elements have comparable diffusivity. If all alloying elements are substitutional but they have remarkable diffusivity difference, e.g. in Al-Zr system, or if the diffusivity is strongly composition-dependent, the *General model* is preferred. If the supersaturation is high, and meanwhile there are fast-diffusing interstitial elements such as C, the *Advanced model* is more appropriate to capture the NPLE mechanism.

# ADVANCED = 3

The **advanced model** has been proposed by *Chen, Jeppsson, and Ågren (CJA) (2008)* and calculates the velocity of a moving phase interface in multicomponent systems by identifying the operating tie-line from the solution of the flux-balance equations. This model can treat both high supersaturation and cross-diffusion rigorously. Spontaneous transitions between different modes (LE and NPLE) of phase transformation can be captured without any ad-hoc treatment.

**Note:** Since it is not always possible to solve the flux-balance equations and it takes time, usage of a less rigorous but simple and efficient model is preferred if possible.

# GENERAL = 5

The **general model** is based on the *Morral-Purdy* model, which follows the same quasi-steady state approximation as the *Simplified model*, but improves it by taking the cross-diffusion into account.

# NPLE = 11

The **Non-Partitioning Local Equilibrium (NPLE) growth rate model** is only available for alloy systems where *Fe* is the major element and at least one interstitial element partitions into the precipitate phase. *This model is specifically designed to deal with the fast diffusion of interstitial elements (C, N, etc.) in Fe alloys.* Based on the *Simplified growth model*, it still holds a local equilibrium condition at the migrating interface. It chooses a tie-line under NPLE condition so that the u-fractions of all substitutional elements and minor

interstitial elements in the precipitate phase are the same as those in the far-field matrix phase (i.e. the overall instantaneous matrix composition).

## $PARA_EQ = 10$

The **para-equilibrium model** is only available for alloy systems where Fe is the major element and C is *the only interstitial element*, which also partitions into the precipitate phase. The interstitial elements, e.g. C, N, etc., usually have remarkably faster diffusion rate than the substitutional elements. Meanwhile, they are assumed to have negligible volume contribution, and as a result the composition variables are replaced by u-fractions when interstitial elements are included in the system. This model is specifically designed to address the fast diffusion of C in Fe alloys. Based on the Simplified growth rate model it holds a para-equilibrium condition at the migrating interface. Contrary to the regular ortho-equilibrium condition state that assumes that all alloying elements are in equilibrium at the interface, the para-equilibrium assumes only equilibrium for C. The substitutional elements are immobile and thus have the same compositions (u-fractions) across the interface.

# $PE\_AUTOMATIC = 12$

The PE Automatic model enables the smooth transition from Paraequilibrium growth rate model to Simplified growth rate model. The rate of transition process is dependent on the relative differences in diffusion between C and substitutional elements, as well as the differences in driving force between paraequilibrium and ortho-equilibrium.

#### SIMPLIFIED = 2

The **simplified model** is based on the *advanced model* but avoids the difficulty of finding the operating tie-line and uses instead the tie-line across the bulk composition. **This is the default growth rate model**.

# class tc\_python.precipitation.MatrixPhase(matrix\_phase\_name: str)

Bases: object

The matrix phase in a precipitation calculation

#### add\_precipitate\_phase(precipitate\_phase: PrecipitatePhase)

Adds a precipitate phase.

```
Parameters
precipitate_phase – The precipitate phase
```

#### 

Enter a numerical value. Default: 5.0E12 m^-2.

#### **Parameters**

dislocation\_density – The dislocation density [m^-2]

**set\_mobility\_adjustment**(*element: str* = '*all*', *prefactor: float* = 1.0, *activation\_energy: float* = 0.0)

A value that adds to the activation energy of mobility data from the database.

#### **Parameters**

- **element** The alement to apply the adjustment for. If "all" is given, adjustment will apply to all elements.
- **prefactor** A parameter that multiplies to the mobility data from a database. This value scales the mobility by a constant amount. This can be useful, for example, when the material has a higher than normal vacancy concentration at the start of the precipitation simulation (e.g. from a prior solutionizing and quenching treatment).
- activation\_energy A value that adds to the activation energy of mobility data from a database. It scales the mobility by a temperature dependent amount. Similar usage as mobility adjustment prefactor. [J/mol]

# set\_molar\_volume(volume: float)

Sets the molar volume of the phase.

**Default**: If not set, the molar volume is taken from the thermodynamic database (or set to 7.0e-6 m<sup>3</sup>/mol if the database contains no molar volume information).

# **Parameters**

volume - The molar volume [m^3/mol]

# with\_elastic\_properties\_cubic(c11: float, c12: float, c44: float)

Sets the elastic properties to "cubic" and specifies the elastic stiffness tensor components. **Default**: if not chosen, the default is DISREGARD

# Parameters

- c11 The stiffness tensor component c11 [GPa]
- c12 The stiffness tensor component c12 [GPa]
- c44 The stiffness tensor component c44 [GPa]

# with\_elastic\_properties\_disregard()

Set to disregard to ignore the elastic properties. Default: This is the default option

# with\_elastic\_properties\_isotropic(shear\_modulus: float, poisson\_ratio: float)

Sets elastic properties to isotropic. Default: if not chosen, the default is DISREGARD

# Parameters

- shear\_modulus The shear modulus [GPa]
- **poisson\_ratio** The Poisson's ratio [-]

# with\_grain\_growth\_model(grain\_growth\_model: GrainGrowthModel)

Sets the model for grain growth. Either fixed size or with a starting distribution

Default: Fixed grain radius size 1.0E-4 m

# **Parameters**

grain\_growth\_model - the grain growth model

# class tc\_python.precipitation.NumericalParameters

Bases: object

Numerical parameters

# set\_max\_overall\_volume\_change(max\_overall\_volume\_change: float = 0.001)

This defines the maximum absolute (not ratio) change of the volume fraction allowed during one time step. **Default**: 0.001

# **Parameters**

**max\_overall\_volume\_change** – The maximum absolute (not ratio) change of the volume fraction allowed during one time step [-]

# set\_max\_radius\_points\_per\_magnitude(max\_radius\_points\_per\_magnitude: float = 200.0)

Sets the maximum number of grid points over one order of magnitude in radius. Default: 200.0

# Parameters

**max\_radius\_points\_per\_magnitude** – The maximum number of grid points over one order of magnitude in radius [-]

set\_max\_rel\_change\_critical\_radius(max\_rel\_change\_critical\_radius: float = 0.1)

Used to place a constraint on how fast the critical radium can vary, and thus put a limit on time step. **Default**: 0.1

Parameters

**max\_rel\_change\_critical\_radius** – The maximum relative change of the critical radius [-]

**set\_max\_rel\_change\_nucleation\_rate\_log**(*max\_rel\_change\_nucleation\_rate\_log: float = 0.5*)

This parameter ensures accuracy for the evolution of effective nucleation rate. Default: 0.5

# **Parameters**

**max\_rel\_change\_nucleation\_rate\_log** – The maximum logarithmic relative change of the nucleation rate [-]

set\_max\_rel\_radius\_change(max\_rel\_radius\_change: float = 0.01)

The maximum value allowed for relative radius change in one time step. Default: 0.01

#### **Parameters**

max\_rel\_radius\_change - The maximum relative radius change in one time step [-]

**set\_max\_rel\_solute\_composition\_change**(*max\_rel\_solute\_composition\_change: float = 0.01*)

Set a limit on the time step by controlling solute depletion or saturation, especially at isothermal stage. **Default**: 0.01

#### Parameters

max\_rel\_solute\_composition\_change - The limit for the relative solute composition
change [-]

# set\_max\_time\_step(max\_time\_step: float = 0.1)

The maximum time step allowed for time integration as fraction of the simulation time. Default: 0.1

#### **Parameters**

max\_time\_step - The maximum time step as fraction of the simulation time [-]

# set\_max\_time\_step\_during\_heating(max\_time\_step\_during\_heating: float = 1.0)

The upper limit of the time step that has been enforced in the heating stages. Default: 1.0 s

#### **Parameters**

**max\_time\_step\_during\_heating** – The maximum time step during heating [s]

0.01)

Sets the maximum volume fraction of subcritical particles allowed to dissolve in one time step. **Default**: 0.01

#### **Parameters**

**max\_volume\_fraction\_dissolve\_time\_step** – The maximum volume fraction of subcritical particles allowed to dissolve in one time step [-]

**set\_min\_radius\_nucleus\_as\_particle**(*min\_radius\_nucleus\_as\_particle: float = 5e-10*)

The cut-off lower limit of precipitate radius. Default: 5.0E-10 m

### **Parameters**

**min\_radius\_nucleus\_as\_particle** – The minimum radius of a nucleus to be considered as a particle [m]

# set\_min\_radius\_points\_per\_magnitude(min\_radius\_points\_per\_magnitude: float = 100.0)

Sets the minimum number of grid points over one order of magnitude in radius. Default: 100.0
#### Parameters

**min\_radius\_points\_per\_magnitude** – The minimum number of grid points over one order of magnitude in radius [-]

# set\_pre\_processing\_option(preprocessing\_option: PreProcessingOption)

Sets the option for enabling pre-processing equilibrium data as a function of temperature. This can be extended to include incipient melting where the incipient melting temperature is approximated for each phase. If the temperature exceeds the incipient melting temperature, the precipitates are removed from the system.

#### **Parameters**

**preprocessing\_option** – Specifies if the preprocessing is either turned off, enabled, or enabled with incipient melting.

set\_radius\_points\_per\_magnitude(radius\_points\_per\_magnitude: float = 150.0)

Sets the number of grid points over one order of magnitude in radius. Default: 150.0

# Parameters

radius\_points\_per\_magnitude - The number of grid points over one order of magnitude
in radius [-]

set\_rel\_radius\_change\_class\_collision(rel\_radius\_change\_class\_collision: float = 0.5)

Sets the relative radius change for avoiding class collision. Default: 0.5

#### **Parameters**

**rel\_radius\_change\_class\_collision** – The relative radius change for avoiding class collision [-]

## class tc\_python.precipitation.ParticleSizeDistribution

Bases: object

Represents the state of a microstructure evolution at a certain time including its particle size distribution, composition and overall phase fraction.

#### add\_radius\_and\_number\_density(radius: float, number\_density: float)

Adds a radius and number density pair to the particle size distribution.

#### Parameters

- radius The radius [m]
- number\_density The number of particles per unit volume per unit length [m^-4]

#### Returns

This ParticleSizeDistribution object

## set\_initial\_composition(element\_name: str, composition\_value: float)

Sets the initial precipitate composition.

# Parameters

- element\_name The name of the element
- **composition\_value** The composition value [composition unit defined for the calculation]

## Returns

This ParticleSizeDistribution object

# set\_volume\_fraction\_of\_phase\_type(volume\_fraction\_of\_phase\_type\_enum: VolumeFractionOfPhaseType)

Sets the type of the phase fraction or percentage. Default: By default volume fraction is used.

Parameters

volume\_fraction\_of\_phase\_type\_enum - Specifies if volume percent or fraction is used

Returns

This ParticleSizeDistribution object

# set\_volume\_fraction\_of\_phase\_value(value: float)

Sets the overall volume fraction of the phase (unit based on the setting of set\_volume\_fraction\_of\_phase\_type()).

Parameters

**value** – The volume fraction 0.0 - 1.0 or percent value 0 - 100

Returns

This ParticleSizeDistribution object

class tc\_python.precipitation.PreProcessingOption(value)

Bases: Enum

Options for preprocessing.

OFF = 0

By default, the equilibrium properties are not characterized prior to a simulation.

ON = 1

The equilibrium properties are characterized as a function of temperature prior to the simulation.

#### ON\_WITH\_INCIPIENT\_MELTING = 2

The equilibrium properties are characterized as a function of temperature, and the approximate incipient melting temperatures are determined for each precipitate phase.

# class tc\_python.precipitation.PrecipitateElasticProperties

Bases: object

Represents the elastic transformation strain of a certain precipitate class.

**Note:** This class is only relevant if the option *TransformationStrainCalculationOption.USER\_DEFINED* has been chosen using *PrecipitatePhase.set\_transformation\_strain\_calculation\_option()*. The elastic strain can only be considered for non-spherical precipitates.

# set\_e11(e11: float)

Sets the elastic strain tensor component e11. Default: 0.0

# Parameters

e11 – The elastic strain tensor component e11

Returns

This PrecipitateElasticProperties object

# set\_e12(e12: float)

Sets the strain tensor component e12. Default: 0.0

## Parameters

e12 – The elastic strain tensor component e12

# Returns

This PrecipitateElasticProperties object

## set\_e13(e13: float)

Sets the elastic strain tensor component e13. Default: 0.0

#### Parameters

e13 – The elastic strain tensor component e13

#### Returns

This PrecipitateElasticProperties object

# set\_e22(e22: float)

Sets the elastic strain tensor component e22. Default: 0.0

# Parameters

e22 – The elastic strain tensor component e22

#### Returns

This PrecipitateElasticProperties object

# set\_e23(e23: float)

Sets the elastic strain tensor component e23. Default: 0.0

# Parameters e23 – The elastic strain tensor component e23

Returns

This PrecipitateElasticProperties object

# set\_e33(e33: float)

Sets the elastic strain tensor component e33. Default: 0.0

# Parameters

e33 – The elastic strain tensor component e33

#### Returns

This PrecipitateElasticProperties object

# class tc\_python.precipitation.PrecipitateMorphology(value)

# Bases: Enum

Available precipitate morphologies.

# CUBOID = 3

Cuboidal precipitates, only available for bulk nucleation.

# NEEDLE = 1

Needle-like precipitates, only available for bulk nucleation.

# PLATE = 2

Plate-like precipitates, only available for bulk nucleation.

# SPHERE = 0

Spherical precipitates, this is the default morphology.

# class tc\_python.precipitation.PrecipitatePhase(precipitate\_phase\_name: str)

Bases: object

Represents a certain precipitate class (i.e. a group of precipitates with the same phase and settings).

# disable\_calculate\_aspect\_ratio\_from\_elastic\_energy()

Disables the automatic calculation of the aspect ratio from the elastic energy of the phase.

This PrecipitatePhase object

**Note:** If you use this method, you are required to set the aspect ratio explicitly using the method set\_aspect\_ratio\_value().

**Default**: This is the default setting (with an aspect ratio of 1.0).

# disable\_driving\_force\_approximation()

Disables driving force approximation for this precipitate class. **Default**: Driving force approximation is disabled.

## Returns

This PrecipitatePhase object

# enable\_calculate\_aspect\_ratio\_from\_elastic\_energy()

Enables the automatic calculation of the aspect ratio from the elastic energy of the phase. **Default**: The aspect ratio is set to a value of 1.0.

#### Returns

This PrecipitatePhase object

# enable\_driving\_force\_approximation()

Enables driving force approximation for this precipitate class. This approximation is often required when simulating precipitation of multiple particles that use the same phase description. E.g. simultaneous precipitation of a Metal-Carbide(MC) and Metal-Nitride(MN) if configured as different composition sets of the same phase FCC\_A1. **Default**: Driving force approximation is disabled.

## Returns

This PrecipitatePhase object

**Tip:** Use this if simulations with several compositions sets of the same phase cause problems.

#### set\_alias(alias: str)

Sets an alias string that can later be used to get values from a calculated result. Typically used when having the same phase for several precipitates, but with different nucleation sites. For example two precipitates of the phase M7C3 with nucleation sites in 'Bulk' and at 'Dislocations'. The alias can be used instead of the phase name when retrieving simulated results.

# Parameters

**alias** – The alias string for this class of precipitates

# Returns

This PrecipitatePhase object

**Note:** Typically used when having using the same precipitate phase, but with different settings in the same calculation.

# set\_aspect\_ratio\_value(aspect\_ratio\_value: float)

Sets the aspect ratio of the phase. **Default**: An aspect ratio of 1.0.

Parameters

aspect\_ratio\_value - The aspect ratio value

This PrecipitatePhase object

**Note:** Only relevant if *disable\_calculate\_aspect\_ratio\_from\_elastic\_energy()* is used (which is the default).

# set\_gibbs\_energy\_addition(gibbs\_energy\_addition: float)

Sets a Gibbs energy addition to the Gibbs energy of the phase. Default: 0,0 J/mol

**Parameters** gibbs\_energy\_addition – The Gibbs energy addition [J/mol]

#### Returns

This PrecipitatePhase object

#### set\_interfacial\_energy(interfacial\_energy: Union[str, float])

Sets the interfacial energy. **Default**: If the interfacial energy is not set, it is automatically calculated using a broken-bond model.

#### Parameters

**interfacial\_energy** – A float for the interfacial energy [J/m<sup>2</sup>] or a string containing a particle size and/or temperature dependent function.

# Returns

This PrecipitatePhase object

**Note:** If this method is not called, the calculated interfacial energy shall be used. The calculation of the interfacial energy using a broken-bond model is based on the assumption of an interface between a bccand a fcc-crystal structure with (110) and (111) lattice planes regardless of the actual phases.

A constant float value can be entered. Alternatively, a string can be given describing a function capturing the desired dependency on size and/or temperature. The size is given in terms of radius "r" with units of meters. Temperature is defined by "T" and has units of Kelvin.

For example, a smooth step function from 0.05 to 0.11 with the transition at 3nm, this is the function: ((0.05+0.03) + 0.03\*erf((r-3E-9)/1e-9)))

Search the Thermo-Calc help for "Operators and Functions" for a list of operators you can use.

# set\_interfacial\_energy\_estimation\_prefactor(interfacial\_energy\_estimation\_prefactor: float)

Sets the interfacial energy prefactor. **Default**: Prefactor of 1.0 (only relevant if the interfacial energy is automatically calculated).

#### Parameters

**interfacial\_energy\_estimation\_prefactor** – The prefactor for the calculated interfacial energy

#### Returns

This PrecipitatePhase object

**Note:** The interfacial energy prefactor is an amplification factor for the automatically calculated interfacial energy. Example: *interfacial\_energy\_estimation\_prefactor* =  $2.5 \Rightarrow 2.5 \approx$  calculated interfacial energy

# set\_molar\_volume(volume: float)

Sets the molar volume of the precipitate phase. **Default**: The molar volume obtained from the database. If no molar volume information is present in the database, a value of 7.0e-6 m^3/mol is used.

#### **Parameters**

volume – The molar volume [m^3/mol]

#### Returns

This PrecipitatePhase object

# set\_nucleation\_at\_dislocations(number\_density=-1)

Activates nucleation at dislocations for this class of precipitates. Calling the method overrides any other nucleation setting for this class of precipitates. **Default**: If not set, by default bulk nucleation is chosen.

#### **Parameters**

**number\_density** – Number density of nucleation sites. If not set, the value is calculated based on the matrix settings (grain size, dislocation density) [m^-3].

# Returns

This PrecipitatePhase object

# set\_nucleation\_at\_grain\_boundaries(wetting\_angle: float = 90.0, number\_density: float = -1)

Activates nucleation at grain boundaries for this class of precipitates. Calling the method overrides any other nucleation setting for this class of precipitates. **Default**: If not set, by default bulk nucleation is chosen.

# **Parameters**

- wetting\_angle If not set, a default value of 90 degrees is used
- **number\_density** Number density of nucleation sites. If not set, the value is calculated based on the matrix settings (grain size) [m<sup>-3</sup>].

## Returns

This PrecipitatePhase object

# **set\_nucleation\_at\_grain\_corners**(*wetting\_angle: float = 90, number\_density: float = -1*)

Activates nucleation at grain corners for this class of precipitates. Calling the method overrides any other nucleation setting for this class of precipitates. **Default**: If not set, by default bulk nucleation is chosen.

## Parameters

- wetting\_angle If not set, a default value of 90 degrees is used]
- **number\_density** Number density of nucleation sites. If not set, the value is calculated based on the matrix settings (grain size) [m^-3].

# Returns

This PrecipitatePhase object

# set\_nucleation\_at\_grain\_edges(wetting\_angle=90, number\_density=-1)

Activates nucleation at the grain edges for this class of precipitates. Calling the method overrides any other nucleation setting for this class of precipitates. **Default**: If not set, by default bulk nucleation is chosen.

# **Parameters**

- wetting\_angle If not set, a default value of 90 degrees is used
- **number\_density** Number density of nucleation sites. If not set, the value is calculated based on the matrix settings (grain size) [m<sup>-3</sup>].

# Returns

This *PrecipitatePhase* object

# set\_nucleation\_in\_bulk(number\_density: float = -1.0)

Activates nucleation in the bulk for this class of precipitates. Calling the method overrides any other nucleation setting for this class of precipitates. **Default**: This is the default setting (with an automatically calculated number density).

# Parameters

**number\_density** – Number density of nucleation sites. If not set, the value is calculated based on the matrix settings (molar volume)  $[m^{-3}]$ 

# Returns

This PrecipitatePhase object

# set\_phase\_boundary\_mobility(phase\_boundary\_mobility: float)

Sets the phase boundary mobility. **Default**: 10.0 m<sup>4</sup>/(Js).

# **Parameters**

phase\_boundary\_mobility – The phase boundary mobility [m^4/(Js)]

# Returns

This PrecipitatePhase object

set\_precipitate\_morphology(precipitate\_morphology\_enum: PrecipitateMorphology)

Sets the precipitate morphology. Default: PrecipitateMorphology. SPHERE

# Parameters

precipitate\_morphology\_enum - The precipitate morphology

# Returns

This PrecipitatePhase object

# set\_trans\_interface\_mobility\_adjustment(element: str = 'all', prefactor: float = 1.0,

#### activation\_energy: float = 0.0)

Trans-interface mobility adjustment Only relevant when growth rate model is PE Automatic A value that adds to the activation energy of mobility data from the database.

# **Parameters**

- **element** The element to apply the adjustment for. If "all" is given, adjustment will apply to all elements.
- **prefactor** A parameter that multiplies to the mobility data from a database. The value scales the mobility by a constant amount. This results in the trans-interface mobility that controls the kinetics of Para-Equilibrium to Ortho-Equilibrium transition.
- **activation\_energy** A value that adds to the activation energy of mobility data from a database. It scales the mobility by a temperature dependent amount. Similar usage as trans-interface mobility adjustment prefactor. [J/mol]

# set\_transformation\_strain\_calculation\_option(transformation\_strain\_calculation\_option\_enum: TransformationStrainCalculationOption)

Sets the transformation strain calculation option. **Default**: *TransformationStrainCalculationOption*. *DISREGARD*.

# Parameters

transformation\_strain\_calculation\_option\_enum - The chosen option

# Returns

This PrecipitatePhase object

set\_zener\_pinning\_parameters(cutoff\_size: float = 7e-07, kinetic\_prefactor: float = 0.5, exponent: float = 0.93)

These parameters are only relevant when zener pinning is enabled in the matrix phase

# **Parameters**

- **cutoff\_size** Precipitates with radius smaller than this value are neglected in pinning force calculation.
- kinetic\_prefactor Dimensionless kinetic coefficient in Zener equation.
- exponent Exponent of precipitate volume fraction in Zener equation.

#### Returns

This PrecipitatePhase object

## with\_elastic\_properties(elastic\_properties: PrecipitateElasticProperties)

Sets the elastic properties. Default: The elastic transformation strain is disregarded by default.

#### Parameters

elastic\_properties – The elastic properties object

#### Returns

This PrecipitatePhase object

**Note:** This method has only an effect if the option *TransformationStrainCalculationOption*. *USER\_DEFINED* is chosen using the method *set\_transformation\_strain\_calculation\_option()*.

# with\_growth\_rate\_model(growth\_rate\_model\_enum: GrowthRateModel)

Sets the growth rate model for the class of precipitates. Default: GrowthRateModel.SIMPLIFIED

# Parameters growth\_rate\_model\_enum – The growth rate model

Returns

This PrecipitatePhase object

# with\_particle\_size\_distribution(particle\_size\_distribution: ParticleSizeDistribution)

Sets the initial particle size distribution for this class of precipitates. **Default**: If the initial particle size distribution is not explicitly provided, the simulation will start from a supersaturated matrix.

# **Parameters**

particle\_size\_distribution – The initial particle size distribution object

#### Returns

This PrecipitatePhase object

Tip: Use this option if you want to study the further evolution of an existing microstructure.

# class tc\_python.precipitation.PrecipitationCCTCalculation(calculation)

Bases: AbstractCalculation

Configuration for a Continuous-Cooling-Time (CCT) precipitation calculation.

# calculate(timeout\_in\_minutes: float = 0.0) $\rightarrow$ PrecipitationCalculationTTTorCCTResult

Runs the CCT diagram calculation.

# Parameters

**timeout\_in\_minutes** – Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a Unre-coverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

#### Returns

A *PrecipitationCalculationTTTorCCTResult* which later can be used to get specific values from the calculated result

# $get_system_data() \rightarrow SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using *with\_system\_modifications()*.

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as \*.tdb-file.

#### Returns

The system data

# set\_composition(element\_name: str, value: float)

Sets the composition of the elements. The unit for the composition can be changed using set\_composition\_unit(). Default: Mole percent (CompositionUnit.MOLE\_PERCENT)

#### **Parameters**

• **element\_name** – The element

• value - The composition (fraction or percent depending on the composition unit)

#### Returns

This PrecipitationCCTCalculation object

#### set\_composition\_unit(unit\_enum: CompositionUnit)

Sets the composition unit. Default: Mole percent (CompositionUnit.MOLE\_PERCENT).

#### Parameters

unit\_enum – The new composition unit

#### Returns

This PrecipitationCCTCalculation object

# set\_cooling\_rates(cooling\_rates: List[float])

Sets all cooling rates for which the CCT diagram should be calculated.

#### Parameters

**cooling\_rates** – A list of cooling rates [K/s]

#### Returns

This PrecipitationCCTCalculation object

# set\_max\_temperature(max\_temperature: float)

Sets maximum temperature of the CCT diagram.

# Parameters

**max\_temperature** – the maximum temperature [K]

# Returns

This PrecipitationCCTCalculation object

#### set\_min\_temperature(min\_temperature: float)

Sets the minimum temperature of the CCT diagram.

# Parameters

**min\_temperature** – the minimum temperature [K]

#### Returns

This PrecipitationCCTCalculation object

# stop\_at\_volume\_fraction\_of\_phase(stop\_criterion\_value: float)

Sets the stop criterion as a volume fraction of the phase. This setting is applied to all phases.

# **Parameters**

**stop\_criterion\_value** – the volume fraction of the phase (a value between 0 and 1)

#### Returns

This PrecipitationCCTCalculation object

# with\_matrix\_phase(matrix\_phase: MatrixPhase)

Sets the matrix phase.

# Parameters matrix\_phase – The matrix phase

Returns

This PrecipitationCCTCalculation object

# with\_numerical\_parameters(numerical\_parameters: NumericalParameters)

Sets the numerical parameters. If not specified, reasonable defaults are be used.

# **Parameters**

numerical\_parameters – The parameters

#### Returns

This PrecipitationCCTCalculation object

# with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

**Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

#### **Parameters**

system\_modifications – The system modification to be performed

# Returns

This PrecipitationCCTCalculation object

## class tc\_python.precipitation.PrecipitationCalculationResult(result)

# Bases: AbstractResult

Result of a precipitation calculation. This can be used to query for specific values.

# save\_to\_disk(path: str)

Saves the result to disc. Note that a result is a folder, containing potentially many files. The result can later be loaded with load\_result\_from\_disk()

# Parameters

**path** – the path to the folder you want the result to be saved in. It can be relative or absolute.

#### Returns

this PrecipitationCalculationResult object

# class tc\_python.precipitation.PrecipitationCalculationSingleResult(result)

Bases: PrecipitationCalculationResult

Result of a isothermal or non-isothermal precipitation calculation. This can be used to query for specific values.

Search the Thermo-Calc help for definitions of the axis variables, e.g. search *isothermal variables* or *non-isothermal variables*.

get\_aspect\_ratio\_distribution\_for\_particle\_length\_of(precipitate\_id: str, time: float) → [List[float], List[float]]

Returns the aspect ratio distribution of a precipitate in dependency of its mean particle length at a certain time.

Only available if the morphology is set to *PrecipitateMorphology.NEEDLE* or *PrecipitateMorphology.PLATE*.

#### **Parameters**

• time – The time [s]

• precipitate\_id – The id of a precipitate can either be the phase name or an alias

#### Returns

A tuple of two lists of floats (mean particle length [m], aspect ratio)

# get\_aspect\_ratio\_distribution\_for\_radius\_of(precipitate\_id: str, time: float) $\rightarrow$ [List[float], List[float]]

Returns the aspect ratio distribution of a precipitate in dependency of its mean radius at a certain time.

Only available if the morphology is set to *PrecipitateMorphology.NEEDLE* or *PrecipitateMorphology.PLATE*.

# Parameters

• time – The time [s]

• precipitate\_id – The id of a precipitate can either be the phase name or an alias

#### Returns

A tuple of two lists of floats (mean radius [m], aspect ratio)

get\_critical\_radius\_of(precipitate\_id: str) → [List[float], List[float]]

Returns the critical radius of a precipitate in dependency of the time.

#### **Parameters**

precipitate\_id – The id of a precipitate can either be phase name or alias

#### Returns

A tuple of two lists of floats (time [s], critical radius [m])

# get\_cubic\_factor\_distribution\_for\_particle\_length\_of(precipitate\_id: str, time: float) →

[List[float], List[float]]

Returns the cubic factor distribution of a precipitate in dependency of its mean particle length at a certain time.

Only available if the morphology is set to *PrecipitateMorphology.CUBOID*.

**Parameters** 

- time The time in seconds
- precipitate\_id The id of a precipitate can either be the phase name or an alias

A tuple of two lists of floats (particle length [m], cubic factor)

 $get\_cubic\_factor\_distribution\_for\_radius\_of(precipitate\_id: str, time: float) \rightarrow [List[float]], \\List[float]]$ 

Returns the cubic factor distribution of a precipitate in dependency of its mean radius at a certain time. Only available if the morphology is set to *PrecipitateMorphology.CUBOID*.

# **Parameters**

• time – The time [s]

• precipitate\_id – The id of a precipitate can either be the phase name or an alias

#### Returns

A tuple of two lists of floats (radius [m], cubic factor)

# get\_driving\_force\_of(precipitate\_id: str) → [List[float], List[float]]

Returns the (by R \* T) normalized driving force of a precipitate in dependency of the time.

#### Parameters

precipitate\_id - The id of a precipitate can either be the phase name or an alias

#### Returns

A tuple of two lists of floats (time [s], normalized driving force)

# $\texttt{get\_grain\_critical\_radius()} \rightarrow [\texttt{List[float]}, \texttt{List[float]}]$

Returns the critical radius of grains in dependency of the time.

#### Returns

A tuple of two lists of floats (time [s], critical radius [m])

# $\texttt{get\_grain\_mean\_radius()} \rightarrow [\texttt{List[float]}, \texttt{List[float]}]$

Returns the mean grain size of the matrix phase in dependency of the time.

#### Returns

A tuple of two lists of floats (time [s], mean radius [m])

#### **get\_grain\_number\_density**() → [List[float], List[float]]

Returns the grain number density (concentration) in dependency of the time.

#### Returns

A tuple of two lists of floats (time [s], grain number density [m^-3])

# get\_grain\_number\_density\_distribution\_for\_length(time: float) → [List[float], List[float]]

Returns the number density distribution of grains in terms of length at the requested time(s).

# **Parameters**

**time** – The time [s]

#### Returns

A tuple of two lists of floats (grain length[m], number of grains per unit volume in a size class [m^-3])

# get\_grain\_number\_density\_distribution\_for\_radius(*time: float*) → [List[float], List[float]]

Returns the number density distribution of a grains in terms of grain radius at the requested time(s).

#### **Parameters**

time – The time [s]

A tuple of two lists of floats (radius [m], number of grains per unit volume in a size class  $[m^{-3}]$ )

# **get\_grain\_size\_distribution**(*time: float*) → [List[float], List[float]]

Returns the size distribution of the matrix phase in dependency of its grain radius at the requested time(s).

# Parameters

time – The time [s]

#### Returns

A tuple of two lists of floats (grain radius[m], number density of grains[m^-4])

# get\_matrix\_composition\_in\_mole\_fraction\_of(element\_name: str) o [List[float], List[float]]

Returns the matrix composition (as mole fractions) of a certain element in dependency of the time.

# Parameters

element\_name - The element

#### Returns

A tuple of two lists of floats (time [s], mole fraction)

# $\texttt{get\_matrix\_composition\_in\_weight\_fraction\_of(\textit{element\_name: str}) \rightarrow [\texttt{List[float]},\texttt{List[float]}]$

Returns the matrix composition (as weight fraction) of a certain element in dependency of the time.

#### **Parameters**

element\_name - The element

#### Returns

A tuple of two lists of floats (time [s], weight fraction)

## **get\_mean\_aspect\_ratio\_of**(*precipitate\_id: str*) → [List[float], List[float]]

Returns the mean aspect ratio of a precipitate in dependency of the time.

Only available if the morphology is set to *PrecipitateMorphology.NEEDLE* or *PrecipitateMorphology.PLATE*.

#### Parameters

precipitate\_id - The id of a precipitate can either be the phase name or an alias

#### Returns

A tuple of two lists of floats (time [s], mean aspect ratio)

# **get\_mean\_cubic\_factor\_of**(*precipitate\_id: str*) → [List[float], List[float]]

Returns the mean cubic factor of a precipitate in dependency of the time. Only available if the morphology is set to *PrecipitateMorphology.CUBOID*.

# Parameters

precipitate\_id - The id of a precipitate can either be the phase name or an alias

#### Returns

A tuple of two lists of floats (time [s], mean cubic factor)

# $\texttt{get\_mean\_particle\_length\_of}(\textit{precipitate\_id: str}) \rightarrow [\texttt{List[float]},\texttt{List[float]}]$

Returns the mean particle length of a precipitate in dependency of the time.

Only available if the morphology is set to *PrecipitateMorphology.NEEDLE* or *PrecipitateMorphology.PLATE*.

# Parameters

precipitate\_id - The id of a precipitate can either be the phase name or an alias

A tuple of two lists of floats (time [s], mean particle length [m])

# **get\_mean\_radius\_2d\_of**(*precipitate\_id: str*) → [List[float], List[float]]

Returns the mean radius of cross-sections taken through the dispersion in dependency of the time.

# Parameters

precipitate\_id – The id of a precipitate can either be phase name or alias

#### Returns

A tuple of two lists of floats (time [s], mean radius [m])

#### **get\_mean\_radius\_of**(*precipitate\_id: str*) → [List[float], List[float]]

Returns the mean radius of a precipitate in dependency of the time.

#### Parameters

precipitate\_id - The id of a precipitate can either be phase name or alias

#### Returns

A tuple of two lists of floats (time [s], mean radius [m])

## get\_normalized\_grain\_size\_distribution(time: float) → [List[float], List[float]]

Returns the normalized number density distribution with the grain radius normalized by the mean radius, for the requested time(s).

#### **Parameters**

**time** – The time [s]

#### Returns

A tuple of two lists of floats (Normalized size, Probability)

 $\label{eq:get_normalized_number_density_distribution_2d_of(\textit{precipitate_id: str, time: float}) \rightarrow [List[float]], \\ List[float]]$ 

Returns the normalized number density distribution of a precipitate in terms of the radius of cross-sections created by taking a plane through the dispersion, normalized by the mean radius of the cross-section, for the requested time(s).

# Parameters

- time The time [s]
- precipitate\_id The id of a precipitate can either be the phase name or an alias

#### Returns

A tuple of two lists of floats (Normalized size, Probability)

 $\label{eq:get_normalized_number_density_distribution_of(\textit{precipitate_id: str, time: float}) \rightarrow [List[float], \\ List[float]]$ 

Returns the normalized number density distribution with the particle radius normalized by the mean radius, for the requested time(s).

# Parameters

- time The time [s]
- precipitate\_id The id of a precipitate can either be the phase name or an alias

# Returns

A tuple of two lists of floats (Normalized size, Probability)

# **get\_nucleation\_rate\_of**(*precipitate\_id: str*) → [List[float], List[float]]

Returns the nucleation rate of a precipitate in dependency of the time.

# Parameters

precipitate\_id – The id of a precipitate can either be the phase name or an alias

#### Returns

A tuple of two lists of floats (time [s], nucleation rate [m^-3 s^-1)

 $\label{eq:get_number_density_distribution_2d_for_particle_length_of(\textit{precipitate_id: str, time: float}) \rightarrow \\ [List[float], List[float]]$ 

Returns the number density distribution of a precipitate considering the radius of cross-sections created by taking a plane through the dispersion, approximating the particles as spherical, for the requested time(s).

# Parameters

- time The time [s]
- precipitate\_id The id of a precipitate can either be the phase name or an alias

# Returns

A tuple of two lists of floats (particle length[m], number of particles per unit area within a size class  $[m^{-2}]$ )

# $get\_number\_density\_distribution\_2d\_for\_radius\_of(precipitate\_id: str, time: float) \rightarrow [List[float]]$ List[float]]

Returns the number density distribution of a precipitate considering the radius of cross-sections created by taking a plane through the dispersion for the requested time(s).

#### Parameters

• time – The time [s]

• precipitate\_id – The id of a precipitate can either be the phase name or an alias

#### Returns

A tuple of two lists of floats (radius [m], number of particles per unit area within a size class [m^-2])

 $\label{eq:get_number_density_distribution_for_particle_length_of(\textit{precipitate_id: str, time: float}) \rightarrow \\ [List[float], List[float]]$ 

Returns the number density distribution of a precipitate in terms of length for the requested time(s).

#### **Parameters**

- time The time [s]
- precipitate\_id The id of a precipitate can either be the phase name or an alias

# Returns

A tuple of two lists of floats (particle length[m], number of particles in the size class per unit volume  $[m^{-3}]$ )

# 

Returns the number density distribution of a precipitate in terms of radius for the requested time(s).

# Parameters

- time The time [s]
- precipitate\_id The id of a precipitate can either be the phase name or an alias

A tuple of two lists of floats (radius [m], number of particles in the size class per unit volume  $[m^{-3}]$ )

# get\_number\_density\_of(precipitate\_id: str) → [List[float], List[float]]

Returns the particle number density (concentration) of a precipitate in dependency of the time.

# Parameters

precipitate\_id - The id of a precipitate can either be phase name or alias

# Returns

A tuple of two lists of floats (time [s], particle number density [m^-3])

get\_precipitate\_composition\_in\_mole\_fraction\_of(precipitate\_id: str, element\_name: str) → [List[float], List[float]]

Returns the precipitate composition (as mole fractions) of a certain element in dependency of the time.

# Parameters

- precipitate\_id The id of a precipitate can either be phase name or alias
- **element\_name** The element

# Returns

A tuple of two lists of floats (time [s], mole fraction)

 $\label{eq:get_precipitate_composition_in_weight_fraction_of(\textit{precipitate_id: str, element_name: str}) \rightarrow \\ [List[float], List[float]]$ 

Returns the precipitate composition (as weight fraction) of a certain element in dependency of the time.

# **Parameters**

- precipitate\_id The id of a precipitate can either be phase name or alias
- **element\_name** The element

# Returns

A tuple of two lists of floats (time [s], weight fraction)

 $\label{eq:get_size_distribution_2d_for_particle_length_of(\textit{precipitate_id: str, time: float}) \rightarrow [List[float], \\ List[float]]$ 

Returns the size distribution of a precipitate considering the radius of cross-sections created by taking a plane through the dispersion, approximating the particles as spherical, for the requested time(s).

# Parameters

• time – The time [s]

• **precipitate\_id** – The id of a precipitate can either be the phase name or an alias

# Returns

A tuple of two lists of floats (particle length[m], number of particles per unit area per unit length  $[m^{-3}]$ )

 $\texttt{get\_size\_distribution\_2d\_for\_radius\_of}(\textit{precipitate\_id: str, time: float}) \rightarrow [\texttt{List[float]}, \texttt{List[float]}]$ 

Returns the 2d size distribution of a precipitate considering the radius of cross-sections created by taking a plane through the dispersion for the requested time(s).

# Parameters

- time The time [s]
- precipitate\_id The id of a precipitate can either be the phase name or an alias

A tuple of two lists of floats (radius [m], number of particles per unit area per unit length  $[m^{-3}]$ )

 $\texttt{get\_size\_distribution\_for\_particle\_length\_of(\textit{precipitate\_id: str, time: float}) \rightarrow [\texttt{List[float]}, \texttt{float}], \texttt{float}]$ 

List[float]]

Returns the size distribution of a precipitate in terms of length for the requested time(s).

# Parameters

• time – The time [s]

• precipitate\_id – The id of a precipitate can either be the phase name or an alias

# Returns

A tuple of two lists of floats (particle length[m], number of particles per unit volume per unit length  $[m^{-4}]$ )

**get\_size\_distribution\_for\_radius\_of**(*precipitate\_id: str, time: float*)  $\rightarrow$  [List[float], List[float]] Returns the size distribution of a precipitate in terms of radius for the requested time(s).

#### Parameters

• time – The time [s]

• precipitate\_id - The id of a precipitate can either be the phase name or an alias

## Returns

A tuple of two lists of floats (radius [m], number of particles per unit volume per unit length  $[m^{-4}]$ )

# get\_split\_mean\_radius\_of(precipitate\_id: str, isSectioned: bool)

Returns the mean radius of a precipitate in dependency of the time.

# Parameters

precipitate\_id - The id of a precipitate can either be phase name or alias

#### Returns

A tuple of two lists of floats (time [s], mean radius [m])

#### get\_split\_number\_density\_of(precipitate\_id: str, isSectioned: bool)

Returns the mean radius of a precipitate in dependency of the time.

# Parameters

precipitate\_id - The id of a precipitate can either be phase name or alias

#### Returns

A tuple of two lists of floats (time [s], mean radius [m])

# get\_split\_size\_distribution\_for\_radius\_of(precipitate\_id: str, time: float, isSectioned: bool)

Returns the size distribution of a precipitate in terms of radius for the requested time(s).

## **Parameters**

- time The time [s]
- precipitate\_id The id of a precipitate can either be the phase name or an alias

# Returns

A tuple of two lists of floats (radius [m], number of particles per unit volume per unit length [m^-4])

#### **Parameters**

**precipitate\_id** – The id of a precipitate can either be phase name or alias

#### Returns

A tuple of two lists of floats (time [s], mean radius [m])

#### **get\_volume\_fraction\_of**(*precipitate\_id: str*) → [List[float], List[float]]

Returns the volume fraction of a precipitate in dependency of the time.

#### **Parameters**

precipitate\_id – The id of a precipitate can either be the phase name or an alias

#### Returns

A tuple of two lists of floats (time [s], volume fraction)

class tc\_python.precipitation.PrecipitationCalculationTTTorCCTResult(result)

Bases: PrecipitationCalculationResult

Result of a TTT or CCT precipitation calculation.

# get\_result\_for\_precipitate(precipitate\_id: str) → [List[float], List[float]]

Returns the calculated data of a TTT or CCT diagram for a certain precipitate.

# Parameters

precipitate\_id – The id of a precipitate can either be the phase name or an alias

#### Returns

A tuple of two lists of floats (time [s], temp [K])

# class tc\_python.precipitation.PrecipitationIsoThermalCalculation(calculation)

Bases: AbstractCalculation

Configuration for an isothermal precipitation calculation.

#### $calculate(timeout_in_minutes: float = 0.0) \rightarrow PrecipitationCalculationSingleResult$

Runs the isothermal precipitation calculation.

#### Parameters

**timeout\_in\_minutes** – Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a Unre-coverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

#### Returns

A *PrecipitationCalculationSingleResult* which later can be used to get specific values from the calculated result

# $get_system_data() \rightarrow SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using *with\_system\_modifications()*.

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as \*.tdb-file.

#### Returns

The system data

# set\_composition(element\_name: str, value: float)

Sets the composition of the elements. The unit for the composition can be changed using set\_composition\_unit(). Default: Mole percent (CompositionUnit.MOLE\_PERCENT)

**Parameters** 

• element\_name - The element

• value - The composition (fraction or percent depending on the composition unit)

#### Returns

This PrecipitationIsoThermalCalculation object

set\_composition\_unit(unit\_enum: CompositionUnit=CompositionUnit.MOLE\_PERCENT)

Sets the composition unit. **Default**: Mole percent (CompositionUnit.MOLE\_PERCENT).

#### **Parameters**

unit\_enum - The new composition unit

# Returns

This PrecipitationIsoThermalCalculation object

# set\_simulation\_time(simulation\_time: float)

Sets the simulation time.

**Parameters** simulation\_time – The simulation time [s]

# Returns

This PrecipitationIsoThermalCalculation object

#### set\_temperature(temperature: float)

Sets the temperature for the isothermal simulation.

# Parameters

temperature – the temperature [K]

#### Returns

This PrecipitationIsoThermalCalculation object

#### with\_matrix\_phase(matrix\_phase: MatrixPhase)

Sets the matrix phase.

#### Parameters

**matrix\_phase** – The matrix phase

#### Returns

This PrecipitationIsoThermalCalculation object

#### with\_numerical\_parameters(numerical\_parameters: NumericalParameters)

Sets the numerical parameters. If not specified, reasonable defaults are be used.

# Parameters

**numerical\_parameters** – The parameters

#### Returns

This PrecipitationIsoThermalCalculation object

# with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions). **Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

#### **Parameters**

system\_modifications – The system modification to be performed

#### Returns

This PrecipitationIsoThermalCalculation object

class tc\_python.precipitation.PrecipitationNonIsoThermalCalculation(calculation)

Bases: AbstractCalculation

Configuration for a non-isothermal precipitation calculation.

#### **calculate**(*timeout\_in\_minutes:* float = 0.0) $\rightarrow$ *PrecipitationCalculationSingleResult*

Runs the non-isothermal precipitation calculation.

#### **Parameters**

**timeout\_in\_minutes** – Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

#### Returns

A *PrecipitationCalculationSingleResult* which later can be used to get specific values from the calculated result

# $get_system_data() \rightarrow SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with\_system\_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as \*.tdb-file.

#### Returns

The system data

# set\_composition(element\_name: str, value: float)

Sets the composition of the elements. The unit for the composition can be changed using set\_composition\_unit(). Default: Mole percent (CompositionUnit.MOLE\_PERCENT)

#### Parameters

element\_name – The element

• value - The composition (fraction or percent depending on the composition unit)

#### Returns

This PrecipitationIsoThermalCalculation object

#### set\_composition\_unit(unit\_enum: CompositionUnit)

Sets the composition unit. **Default**: Mole percent (CompositionUnit.MOLE\_PERCENT).

#### **Parameters**

unit\_enum – The new composition unit

This PrecipitationIsoThermalCalculation object

# set\_simulation\_time(simulation\_time: float)

Sets the simulation time.

# Parameters

simulation\_time - The simulation time [s]

## Returns

This PrecipitationNonThermalCalculation object

# with\_matrix\_phase(matrix\_phase: MatrixPhase)

Sets the matrix phase.

Parameters matrix\_phase – The matrix phase

# Returns

# This PrecipitationIsoThermalCalculation object

## with\_numerical\_parameters(numerical\_parameters: NumericalParameters)

Sets the numerical parameters. If not specified, reasonable defaults are be used.

# Parameters

numerical\_parameters – The parameters

#### Returns

This PrecipitationIsoThermalCalculation object

# with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

**Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

#### **Parameters**

system\_modifications - The system modification to be performed

## Returns

This PrecipitationNonThermalCalculation object

#### with\_temperature\_profile(temperature\_profile: TemperatureProfile)

Sets the temperature profile to use with this calculation.

# Parameters

**temperature\_profile** – the temperature profile object (specifying time / temperature points)

#### Returns

This PrecipitationNonThermalCalculation object

# class tc\_python.precipitation.PrecipitationTTTCalculation(calculation)

# Bases: AbstractCalculation

Configuration for a TTT (Time-Temperature-Transformation) precipitation calculation.

**calculate**(*timeout\_in\_minutes: float* = 0.0)  $\rightarrow$  *PrecipitationCalculationTTTorCCTResult* 

Runs the TTT diagram calculation.

#### **Parameters**

**timeout\_in\_minutes** – Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a Unre-coverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

#### Returns

A *PrecipitationCalculationTTTorCCTResult* which later can be used to get specific values from the calculated result.

# $get_system_data() \rightarrow SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with\_system\_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as \*.tdb-file.

#### Returns

The system data

# set\_composition(element\_name: str, value: float)

Sets the composition of the elements. The unit for the composition can be changed using set\_composition\_unit(). Default: Mole percent (CompositionUnit.MOLE\_PERCENT)

#### **Parameters**

• element\_name - The element

• value - The composition (fraction or percent depending on the composition unit)

Returns

This PrecipitationTTTCalculation object

# set\_composition\_unit(unit\_enum: CompositionUnit)

Sets the composition unit. Default: Mole percent (CompositionUnit.MOLE\_PERCENT).

#### Parameters

unit\_enum – The new composition unit

#### Returns

This PrecipitationTTTCalculation object

# set\_max\_annealing\_time(max\_annealing\_time: float)

Sets the maximum annealing time, i.e. the maximum time of the simulation if the stopping criterion is not reached.

# **Parameters**

**max\_annealing\_time** – the maximum annealing time [s]

# Returns

This PrecipitationTTTCalculation object

#### set\_max\_temperature(max\_temperature: float)

Sets the maximum temperature for the TTT diagram.

#### **Parameters**

**max\_temperature** – the maximum temperature [K]

This PrecipitationTTTCalculation object

#### set\_min\_temperature(min\_temperature: float)

Sets the minimum temperature for the TTT diagram.

# Parameters

**min\_temperature** – the minimum temperature [K]

#### Returns

This PrecipitationTTTCalculation object

# set\_temperature\_step: float)

Sets the temperature step for the TTT diagram. If not set, the default value is 10 K.

# Parameters temperature\_step – the temperature step [K]

Returns

This PrecipitationTTTCalculation object

# stop\_at\_percent\_of\_equilibrium\_fraction(percentage: float)

Sets the stop criterion to a percentage of the overall equilibrium phase fraction, alternatively a required volume fraction can be specified (using *stop\_at\_volume\_fraction\_of\_phase()*).

#### Parameters

**percentage** – the percentage to stop at (value between 0 and 100)

#### Returns

This PrecipitationTTTCalculation object

## stop\_at\_volume\_fraction\_of\_phase(volume\_fraction: float)

Sets the stop criterion as a volume fraction of the phase, alternatively a required percentage of the equilibrium phase fraction can be specified (using stop\_at\_percent\_of\_equilibria\_fraction()). Stopping at a specified volume fraction is the default setting.

This setting is applied to all phases.

# **Parameters**

**volume\_fraction** – the volume fraction to stop at (a value between 0 and 1)

#### Returns

This PrecipitationTTTCalculation object

#### with\_matrix\_phase(matrix\_phase: MatrixPhase)

Sets the matrix phase.

# Parameters

matrix\_phase - The matrix phase

#### Returns

This PrecipitationTTTCalculation object

# with\_numerical\_parameters(numerical\_parameters: NumericalParameters)

Sets the numerical parameters. If not specified, reasonable defaults are be used.

# Parameters

numerical\_parameters – The parameters

#### Returns

This PrecipitationTTTCalculation object

# with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

**Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

# Parameters

system\_modifications – The system modification to be performed

Returns

This PrecipitationTTTCalculation object

class tc\_python.precipitation.TransformationStrainCalculationOption(value)

Bases: Enum

Options for calculating the transformation strain.

#### CALCULATE\_FROM\_MOLAR\_VOLUME = 2

Calculates the transformation strain from the molar volume, obtains a purely dilatational strain.

#### DISREGARD = 1

Ignores the transformation strain, this is the default setting.

#### $USER_DEFINED = 3$

Transformation strain to be specified by the user.

## class tc\_python.precipitation.VolumeFractionOfPhaseType(value)

Bases: Enum

Unit of the volume fraction of a phase.

# VOLUME\_FRACTION = 6

Volume fraction (0 - 1), this is the default.

#### $VOLUME_PERCENT = 5$

Volume percent (0% - 100%).

# 6.1.4 Module "scheil"

#### class tc\_python.scheil.CalculateSecondaryDendriteArmSpacing

Bases: ScheilBackDiffusion

Configures a secondary dendrite arm spacing calculation used by Scheil *with back diffusion*. The used equation is c \* cooling\_rate^(-n) with c and n being provided either by the user or taken from the defaults.

# disable\_delta\_ferrite\_to\_austenite\_transition()

Turns off the delta ferrite BCC to austenite FCC transition.

**Default**: Delta ferrite to austenite transition is off. :return: This CalculateSecondaryDendriteArmSpacing object

# enable\_delta\_ferrite\_to\_austenite\_transition()

Turns on the delta ferrite BCC to austenite FCC transition.

**Default**: Delta ferrite to austenite transition is off. :return: This CalculateSecondaryDendriteArmSpacing object

#### **set\_c**(*c*: *float* = 5*e*-05)

Sets the scaling factor c in the governing equation c \* cooling\_rate^(-n).

Default: 50 µm

# Parameters

**c** – The scaling factor [m]

#### Returns

This CalculateSecondaryDendriteArmSpacing object

#### set\_cooling\_rate(cooling\_rate: float = 1.0)

Sets the cooling rate.

Default: 1.0 K/s

An increased value moves the result from equilibrium toward a Scheil-Gulliver calculation.

# **Parameters** cooling\_rate – The cooling rate [K/s]

#### Returns

This CalculateSecondaryDendriteArmSpacing object

#### set\_fast\_diffusing\_elements(element\_names: List[str])

Sets elements as fast diffusing. This allows redistribution of these elements in both the solid and liquid parts of the alloy.

Default: No fast-diffusing elements.

# Parameters element\_names – The elements

## Returns

This CalculateSecondaryDendriteArmSpacing object

# **set\_n**(*n*: *float* = 0.33)

Sets the exponent n in the governing equation c \* cooling\_rate^(-n).

#### Default: 0.33

#### **Parameters**

**n** – The exponent [-]

# Returns

This CalculateSecondaryDendriteArmSpacing object

# set\_primary\_phasename(primary\_phase\_name: str = 'AUTOMATIC')

Sets the name of the primary phase.

The primary phase is the phase where the back diffusion takes place. If *AUTOMATIC* is selected, the program tries to find the phase which will give the most back diffusion. That behavior can be overridden by selecting a specific primary phase.

# Default: AUTOMATIC

Parameters

**primary\_phase\_name** – The phase name (or *AUTOMATIC*)

This CalculateSecondaryDendriteArmSpacing object

class tc\_python.scheil.ConstantSecondaryDendriteArmSpacing(secondary\_dendrite\_arm\_spacing:

float = 5e-05)

# Bases: ScheilBackDiffusion

Configures a constant secondary dendrite arm spacing used by Scheil *with back diffusion*. The secondary dendrite arm spacing can either be provided by the user or taken from the defaults.

# disable\_delta\_ferrite\_to\_austenite\_transition()

Turns off the delta ferrite BCC to austenite FCC transition.

**Default**: Delta ferrite to austenite transition is off. :return: This ConstantSecondaryDendriteArmSpacing object

# enable\_delta\_ferrite\_to\_austenite\_transition()

Turns on the delta ferrite BCC to austenite FCC transition.

**Default**: Delta ferrite to austenite transition is off. :return: This ConstantSecondaryDendriteArmSpacing object

#### set\_cooling\_rate(cooling\_rate: float = 1.0)

Sets the cooling rate.

Default: 1.0 K/s

An increased value moves the result from equilibrium toward a Scheil-Gulliver calculation.

Parameters cooling\_rate – The cooling rate [K/s]

#### Returns

This ConstantSecondaryDendriteArmSpacing object

# set\_fast\_diffusing\_elements(element\_names: List[str])

Sets elements as fast diffusing. This allows redistribution of these elements in both the solid and liquid parts of the alloy.

Default: No fast-diffusing elements.

Parameters element\_names – The elements

#### Returns

This ConstantSecondaryDendriteArmSpacing object

# set\_primary\_phasename(primary\_phase\_name: str = 'AUTOMATIC')

Sets the name of the primary phase.

The primary phase is the phase where the back diffusion takes place. If *AUTOMATIC* is selected, the program tries to find the phase which will give the most back diffusion. That behavior can be overridden by selecting a specific primary phase.

#### **Default**: AUTOMATIC

#### Parameters

primary\_phase\_name - The phase name (or AUTOMATIC)

#### Returns

This ConstantSecondaryDendriteArmSpacing object

# class tc\_python.scheil.InterfaceDrivingForceModel(value)

# Bases: Enum

The interface driving force model (free energy change at the liquid/primary solid interface) that is used to evaluate the migration speed in comparison with the maximum velocity.

# DRIVING\_ENERGY = 0

Using the overall free energy change. This is the default.

# $MIGRATION\_ENERGY = 1$

Uses free energy change due to migration, i.e. the overall free energy change excluding that contributes to diffusion.

# class tc\_python.scheil.ScheilBackDiffusion

Bases: ScheilCalculationType

Configuration for back diffusion in the solid primary phase.

**Warning:** This feature has only effect on systems with diffusion data (typically a mobility database). If used for a system without diffusion data, a normal Scheil calculation is done.

# classmethod calculate\_secondary\_dendrite\_arm\_spacing()

Calculate the secondary dendrite arm spacing based on the following equation: c \* cooling\_rate^(-n) with c and n being provided either by the user or taken from the defaults.

Use the methods provide by CalculateSecondaryDendriteArmSpacing to configure the parameters.

#### Returns

A CalculateSecondaryDendriteArmSpacing

Assuming constant secondary dendrite arm spacing, provided either by the user or taken from the defaults.

Default: 50 µm

Parameters secondary\_dendrite\_arm\_spacing – The dendrite arm spacing [m]

# Returns

A ConstantSecondaryDendriteArmSpacing

# class tc\_python.scheil.ScheilCalculation(calculator)

Bases: AbstractCalculation

Configuration for a Scheil solidification calculation.

**Note:** Specify the settings, the calculation is performed with *calculate()*.

# $calculate(timeout_in_minutes: float = 0.0) \rightarrow ScheilCalculationResult$

Runs the Scheil calculation.

# Parameters

**timeout\_in\_minutes** – Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a Unre-coverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

A *ScheilCalculationResult* which later can be used to get specific values from the simulation.

# disable\_global\_minimization()

Disables global minimization.

Default: Enabled

**Note:** When enabled, a global minimization test is performed when an equilibrium is reached. This costs more computer time but the calculations are more robust.

#### Returns

This ScheilCalculation object

# enable\_global\_minimization()

Enables global minimization.

Default: Enabled

**Note:** When enabled, a global minimization test is performed when an equilibrium is reached. This costs more computer time but the calculations are more robust.

#### Returns

This ScheilCalculation object

# $get_system_data() \rightarrow SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with\_system\_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as \*.tdb-file.

#### Returns

The system data

#### set\_composition(component\_name: str, value: float)

Sets the composition of a component. The unit for the composition can be changed using set\_composition\_unit().

**Default**: Mole percent (CompositionUnit.MOLE\_PERCENT)

Parameters

- component\_name The component
- value The composition value [composition unit defined for the calculation]

## Returns

This ScheilCalculation object

set\_composition\_unit(unit\_enum: CompositionUnit = CompositionUnit.MOLE\_PERCENT)
Sets the composition unit.

Default: Mole percent (CompositionUnit.MOLE\_PERCENT).

Parameters

unit\_enum - The new composition unit

Returns

This ScheilCalculation object

set\_start\_temperature(temperature\_in\_kelvin: float = 2500.0)

Sets the start temperature.

Warning: The start temperature needs to be higher than the liquidus temperature of the alloy.

Default: 2500.0 K

**Parameters** temperature\_in\_kelvin – The temperature [K]

Returns

This ScheilCalculation object

with\_calculation\_type(scheil\_calculation\_type: ScheilCalculationType)

Chooses a specific Scheil calculation.

- ClassicScheil for only setting fast diffusers.
- *ScheilBackDiffusion* enables back diffusion in the solid primary phase and optionally fast diffusers in all solid phases.
- ScheilSoluteTrapping enables solute trapping in the solid primary phase.

# Parameters

**scheil\_calculation\_type** – Type of Scheil calculation, either *ScheilClassic*, *ScheilBack-Diffusion* or *ScheilSoluteTrapping* 

# Returns

This ScheilCalculation object

# with\_options(options: ScheilOptions)

Sets the Scheil simulation options.

# Parameters

options – The Scheil simulation options

#### Returns

This ScheilCalculation object

## with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

**Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

#### **Parameters**

system\_modifications – The system modification to be performed

#### Returns

This ScheilCalculation object

## class tc\_python.scheil.ScheilCalculationResult(result)

Bases: AbstractResult

Result of a Scheil calculation.

# $\texttt{get\_solid\_phase\_with\_largest\_mole\_fraction()} \rightarrow str$

Returns the name of the solid phase with the largest amount in terms of mole fraction at the end of the Scheil simulation.

#### Returns

Phase name

# $\texttt{get\_stable\_phases()} \rightarrow List[str]$

Returns all phases that were stable during a Scheil simulation.

# Returns

The list of stable phases

# get\_values\_grouped\_by\_quantity\_of(x\_quantity: Union[ScheilQuantity, str], y\_quantity:

*Union*[ScheilQuantity, *str*], *sort\_and\_merge: bool* = *True*) → Dict[str, *ResultValueGroup*]

Returns x-y-line data grouped by the multiple datasets of the specified quantities (for example in dependency of phases or components). Use *get\_values\_of()* instead if you need no separation. The available quantities can be found in the documentation of the factory class ScheilQuantity.

**Note:** The different datasets might contain *NaN*-values between different subsections and might not be sorted **even if the flag `sort\_and\_merge` has been set** (because they might be unsortable due to their nature).

# **Parameters**

- **x\_quantity** The first Scheil quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example "T")
- **y\_quantity** The second Scheil quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example "NV")
- **sort\_and\_merge** If *True*, the data is sorted and merged into as few subsections as possible (divided by *NaN*)

# Returns

Containing the ResultValueGroup dataset objects with their quantity labels as keys

# get\_values\_grouped\_by\_stable\_phases\_of(x\_quantity: Union[ScheilQuantity, str], y\_quantity:

*Union*[ScheilQuantity, *str*], *sort\_and\_merge: bool* = *True*) → Dict[str, *ResultValueGroup*]

Returns x-y-line data grouped by the sets of "stable phases" (for example "LIQUID" or "LIQUID + FCC\_A1"). Use *get\_values\_of()* instead if you need no separation. The available quantities can be found in the documentation of the factory class ScheilQuantity.

**Note:** The different datasets might contain *NaN*-values between different subsections and might not be sorted **even if the flag `sort\_and\_merge` has been set** (because they might be unsortable due to their nature).

# Parameters

- **x\_quantity** The first Scheil quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example "T")
- **y\_quantity** The second Scheil quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example "NV")
- **sort\_and\_merge** If *True*, the data will be sorted and merged into as few subsections as possible (divided by *NaN*)

# Returns

Containing the ResultValueGroup dataset objects with their "stable phases" labels as keys

```
get_values_of(x_quantity: Union[ScheilQuantity, str], y_quantity: Union[ScheilQuantity, str]) \rightarrow
[List[float], List[float]]
```

Returns sorted x-y-line data without any separation. Use get\_values\_grouped\_by\_quantity\_of() or get\_values\_grouped\_by\_stable\_phases\_of() instead if you need such a separation. The available quantities can be found in the documentation of the factory class ScheilQuantity.

**Note:** This method will always return sorted data without any *NaN*-values. In case of ambiguous quantities (for example: CompositionOfPhaseAsWeightFraction("FCC\_A1", "All")) that can give data that is hard to interpret. In such a case you need to choose the quantity in another way or use one of the other methods.

## **Parameters**

- **x\_quantity** The first Scheil quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example "T")
- **y\_quantity** The second Scheil quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example "NV")

# Returns

A tuple containing the x- and y-data in lists

# save\_to\_disk(path: str)

Saves the result to disc. Note that a result is a folder, containing potentially many files. The result can later be loaded with load\_result\_from\_disk()

# Parameters

**path** – the path to the folder you want the result to be saved in.

# Returns

this ScheilCalculationResult object

# class tc\_python.scheil.ScheilCalculationType

Bases: object

Specific configuration for the different Scheil calculation types

## classmethod scheil\_back\_diffusion()

Configuration for back diffusion in the solid primary phase.

**Warning:** This feature has only effect on systems with diffusion data (typically a mobility database). If used for a system without diffusion data, a normal Scheil calculation is done.

# Returns

A ScheilBackDiffusion

# classmethod scheil\_classic()

Configuration for Classic Scheil with fast diffusers.

# Returns

A ScheilClassic

# classmethod scheil\_solute\_trapping()

Configures the Scheil solute trapping settings. The used solidification speed equation is *Scanning speed* \* *cos(angle)* with *Scanning speed* and *angle* being provided either by the user or taken from the defaults.

# Returns

A ScheilSoluteTrapping

# class tc\_python.scheil.ScheilClassic

Bases: ScheilCalculationType

Configuration for Classic Scheil with fast diffusers.

# disable\_delta\_ferrite\_to\_austenite\_transition()

Turns off the delta ferrite BCC to austenite FCC transition.

Default: Delta ferrite to austenite transition is off. :return: This ScheilClassic object

# enable\_delta\_ferrite\_to\_austenite\_transition()

Turns on the delta ferrite BCC to austenite FCC transition.

Default: Delta ferrite to austenite transition is off. :return: This ScheilClassic object

# set\_fast\_diffusing\_elements(element\_names: List[str])

Sets elements as fast diffusing. This allows redistribution of these elements in both the solid and liquid parts of the alloy.

Default: No fast-diffusing elements.

Parameters element\_names – The elements

#### Returns

This ScheilClassic object

# class tc\_python.scheil.ScheilOptions

Bases: object

Options for the Scheil simulation.

# calculate\_from\_gas()

Calculates the evaporation temperature if a gas phase is selected in the system, and then calculates equilibria in the gas+liquid and liquid regions until liquidus temperature is reached.

Default: Calculation starts from liquidus temperature.

This ScheilOptions object

# calculate\_from\_liquidus()

Solidification calculation starting from the liquidus temperature. Liquid properties between start temperature and liquidus are not obtainable.

Default: Calculation starts from liquidus temperature.

Returns

This ScheilOptions object

# calculate\_from\_start\_temperature()

Calculation of equilibria from start temperature at 50 K intervals until liquidus temperature is reached, using the temperature defined with *ScheilCalculation.set\_start\_temperature()*. This option makes it possible to obtain properties of the liquid phase before the solidification starts.

Default: Calculation starts from liquidus temperature.

#### Returns

This ScheilOptions object

# calculate\_to\_end\_of\_scheil()

Stops the calculation when the Scheil calculation is finished.

Default: Calculation stops when the Scheil calculation is finished.

#### Returns

This ScheilOptions object

calculate\_to\_temperature\_below\_solidus(number\_of\_steps: int = 50, final\_temperature: float =

298.15)

Calculates properties in the solid state, for the phase compositions and fractions at the end of the Scheil calculation.

Default: Calculation stops when the Scheil calculation is finished.

#### Parameters

- **number\_of\_steps** Calculates properties for the given number of temperatures, down to the final temperature.
- **final\_temperature** The final (lowest) temperature where the calculation is performed.

# Returns

This ScheilOptions object

#### disable\_approximate\_driving\_force\_for\_metastable\_phases()

Disables the approximation of the driving force for metastable phases.

Default: Enabled

**Note:** When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it is important that these driving forces are correct, use *disable\_approximate\_driving\_force\_for\_metastable\_pha* to force the calculation to converge for the metastable phases.

#### Returns

This ScheilOptions object

# disable\_control\_step\_size\_during\_minimization()

Disables stepsize control during minimization (non-global).

Default: Enabled

# Returns

This ScheilOptions object

# disable\_equilibrium\_solidification\_calculation()

Skips the property (one axis) diagram calculation of solidification under equilibrium conditions, before the Scheil solidification calculation starts.

In general it is not necessary to perform this calculation.

Default: Disabled. The equilibrium solidification calculation is skipped.

#### Returns

This ScheilOptions object

# disable\_evaporation\_property\_calculation()

Disables calculation of evaporation properties.

**Default**: Disabled. The evaporation properties are not calculated.

#### Returns

This ScheilOptions object

# disable\_force\_positive\_definite\_phase\_hessian()

Disables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

Default: Enabled

#### Returns

This ScheilOptions object

# enable\_approximate\_driving\_force\_for\_metastable\_phases()

Enables the approximation of the driving force for metastable phases.

Default: Enabled

**Note:** When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it is important that these driving forces are correct, use *disable\_approximate\_driving\_force\_for\_metastable\_pha* to force the calculation to converge for the metastable phases.

## Returns

This ScheilOptions object

# enable\_control\_step\_size\_during\_minimization()

Enables stepsize control during normal minimization (non-global).

# Default: Enabled

# Returns

This ScheilOptions object

# enable\_equilibrium\_solidification\_calculation()

Performs a property (one axis) diagram calculation of solidification under equilibrium conditions, before the Scheil solidification calculation starts, in the same way as is typically done in graphical and console mode.

In general it is not necessary to perform this calculation.

Default: Disabled. The equilibrium solidification calculation is skipped.

#### Returns

This ScheilOptions object

# enable\_evaporation\_property\_calculation()

Enables calculation of the properties molar mass of gas, driving force for evaporation and evaporation enthalpy. The calculation requires the gas phase to be selected.

Default: Disabled. The evaporation properties are not calculated.

#### Returns

This ScheilOptions object

# enable\_force\_positive\_definite\_phase\_hessian()

Enables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

#### Default: Enabled

Returns

This ScheilOptions object

# set\_gas\_phase(phase\_name: str = 'GAS')

Sets the phase used as the gas phase.

Default: The phase "GAS".

# **Parameters**

phase\_name - The phase name

#### Returns

This ScheilOptions object

#### set\_global\_minimization\_max\_grid\_points(max\_grid\_points: int = 2000)

Sets the maximum number of grid points in global minimization. **\*\*** Only applicable if global minimization is actually used**\*\***.

Default: 2000 points

# Parameters

max\_grid\_points - The maximum number of grid points

#### Returns

This ScheilOptions object

set\_global\_minimization\_test\_interval(global\_test\_interval: int = 10)

Sets the interval for the global test.

# Default: 10

Parameters global\_test\_interval – The global test interval Returns This ScheilOptions object

set\_liquid\_phase(phase\_name: str = 'LIQUID')

Sets the phase used as the liquid phase.

Default: The phase "LIQUID".

Parameters phase\_name – The phase name

Returns This ScheilOptions object

set\_max\_no\_of\_iterations(max\_no\_of\_iterations: int = 500)

Set the maximum number of iterations.

Default: max. 500 iterations

**Note:** As some models give computation times of more than 1 CPU second/iteration, this number is also used to check the CPU time and the calculation stops if 500 CPU seconds/iterations are used.

Parameters

**max\_no\_of\_iterations** – The max. number of iterations

Returns

This ScheilOptions object

# set\_required\_accuracy(accuracy: float = 1e-06)

Sets the required relative accuracy.

Default: 1.0E-6

**Note:** This is a relative accuracy, and the program requires that the relative difference in each variable must be lower than this value before it has converged. A larger value normally means fewer iterations but less accurate solutions. The value should be at least one order of magnitude larger than the machine precision.

**Parameters** 

accuracy – The required relative accuracy

Returns

This ScheilOptions object

# set\_smallest\_fraction(smallest\_fraction: float = 1e-12)

Sets the smallest fraction for constituents that are unstable.

It is normally only in the gas phase that you can find such low fractions.

The **default value** for the smallest site-fractions is 1E-12 for all phases except for IDEAL phase with one sublattice site (such as the GAS mixture phase in many databases) for which the default value is always as 1E-30.

Parameters

smallest\_fraction – The smallest fraction for constituents that are unstable

Returns

This ScheilOptions object
set\_temperature\_step(temperature\_step\_in\_kelvin: float = 1.0)

Sets the temperature step. Decreasing the temperature step increases the accuracy, but the default value is usually adequate.

**Default step:** 1.0 K

```
Parameters
temperature_step_in_kelvin – The temperature step [K]
```

Returns

This ScheilOptions object

terminate\_on\_fraction\_of\_liquid\_phase(fraction\_to\_terminate\_at: float = 0.01)

Sets the termination condition to a specified remaining fraction of liquid phase.

**Default**: Terminates at 0.01 fraction of liquid phase.

**Note:** Either the termination criterion is set to a temperature or fraction of liquid limit, both together are not possible.

#### **Parameters**

**fraction\_to\_terminate\_at** – the termination fraction of liquid phase (value between 0 and 1)

#### Returns

This ScheilOptions object

# terminate\_on\_temperature(temperature\_in\_kelvin: float)

Sets the termination condition to a specified temperature.

Default: Terminates at 0.01 fraction of liquid phase, i.e. not at a specified temperature.

**Note:** Either the termination criterion is set to a temperature or fraction of liquid limit, both together are not possible.

#### **Parameters**

temperature\_in\_kelvin – the termination temperature [K]

#### Returns

This ScheilOptions object

# class tc\_python.scheil.ScheilSoluteTrapping

Bases: ScheilCalculationType

Configures the Scheil solute trapping settings. The used solidification speed equation is *Scanning speed* \* *cos(angle)* with *Scanning speed* and *angle* being provided either by the user or taken from the defaults.

set\_angle(alpha: float = 45.0)

Sets the transformation angle alpha between the solid/liquid boundary and laser scanning direction.

**Default**: 45.0

Parameters alpha – The transformation angle [degree]

#### Returns

This ScheilSoluteTrapping object

```
set_interface_driving_force_model(model: InterfaceDrivingForceModel =
```

InterfaceDrivingForceModel.DRIVING\_ENERGY)

Sets the interface driving force model (free energy change at the liquid/primary solid interface) that is used to evaluate the migration speed in comparison with the maximum velocity.

It can be the DRIVING\_ENERGY or MIGRATION\_ENERGY model.

**Default**: DRIVING\_ENERGY

# Parameters

model – The interface driving force model

Returns

This ScheilSoluteTrapping object

# set\_maximum\_velocity\_for\_infinite\_driving\_force(maximum\_velocity: float = 2000)

Sets the maximum migration speed of the liquid/primary solid interface when the driving force is infinite.

**Default**: 2000 m/s

Parameters maximum\_velocity – The maximum migration speed [m/s]

#### Returns

This ScheilSoluteTrapping object

set\_primary\_phasename(primary\_phase\_name: str = 'AUTOMATIC')

Sets the name of the primary phase.

The primary phase is the phase where solute trapping takes place. A necessary condition for this phase is that the phase definition contains all of the elements that are chosen in the system. When *AUTOMATIC* is selected, the program tries to find a suitable primary phase that fills this condition.

Default: AUTOMATIC

```
Parameters
primary_phase_name – The phase name (or AUTOMATIC)
```

Returns

This ScheilSoluteTrapping object

set\_scanning\_speed(scanning\_speed: float = 1.0)

Sets the scanning speed.

Default: 1 m/s

**Parameters** scanning\_speed – The scaling factor [m/s]

Returns

This ScheilSoluteTrapping object

set\_solute\_trapping\_model(model: SoluteTrappingModel=SoluteTrappingModel.AZIZ)

Sets the solute trapping model for the Scheil calculation.

The model is used to calculate the relation between migration speed and solute partitioning at the liquid/primary solid interface, developed by either Aziz or Hillert.

Default: AZIZ

Parameters

**model** – The solute trapping model

Returns

This ScheilSoluteTrapping object

set\_trans\_interface\_diffusivity\_for(element: str, pre\_factor: float = 5e-09, activation\_energy: float = 0.0)

Sets the solute diffusivity across the interface between liquid and primary dendrite phase **for a specified element**, defined by an Arrhenius-type equation.

**Default**: 5.0e-9 m\*\*2/s for the pre-factor and 0 J/mol for the activation energy

**Note:** If the trans-interface diffusivity had previously been set for *all* elements, this setting will be completely removed.

#### **Parameters**

- element The element for which the trans-interface diffusivity is to be set
- **pre\_factor** The pre-exponential factor in the Arrhenius equation for the trans-interface diffusivity [m\*\*2/s]
- **activation\_energy** The activation energy in the Arrhenius equation for the transinterface diffusivity [J/mol]

#### Returns

This ScheilSoluteTrapping object

Sets the solute diffusivity across the interface between liquid and primary dendrite phase **for all elements**, defined by an Arrhenius-type equation.

**Default**: 5.0e-9 m\*\*2/s for the pre-factor and 0 J/mol for the activation energy

#### **Parameters**

- **pre\_factor** The pre-exponential factor in the Arrhenius equation for the trans-interface diffusivity [m\*\*2/s].
- **activation\_energy** The activation energy in the Arrhenius equation for the transinterface diffusivity [J/mol].

#### Returns

This ScheilSoluteTrapping object

# class tc\_python.scheil.SoluteTrappingModel(value)

# Bases: Enum

The solute trapping model to calculate the relation between migration speed and solute partitioning at the liquid/primary solid interface.

# AZIZ = 0

The solute trapping model developed by Aziz. This is the default.

# HILLERT = 1

The solute trapping model developed by Hillert model.

# 6.1.5 Module "step\_or\_map\_diagrams"

# class tc\_python.step\_or\_map\_diagrams.AbstractAxisType

Bases: object

The abstract base class for all axis types.

# class tc\_python.step\_or\_map\_diagrams.AbstractPhaseDiagramCalculation(calculator)

Bases: AbstractCalculation

Abstract configuration required for a property diagram calculation.

Note: This is an abstract class that cannot be used directly.

#### add\_initial\_equilibrium(initial\_equilibrium: InitialEquilibrium)

Add initial equilibrium start points from which a phase diagram is calculated.

Scans along the axis variables and generates start points when the scan procedure crosses a phase boundary.

It may take a little longer to execute than using the minimum number of start points, as some lines may be calculated more than once. But the core remembers all node points and subsequently stops calculations along a line when it finds a known node point.

It is also possible to create a sequence of start points from one initial equilibria.

#### **Parameters**

initial\_equilibrium – The initial equilibrium

#### Returns

This PhaseDiagramCalculation object

**abstract calculate**(keep\_previous\_results: bool = False, timeout\_in\_minutes: float = 0.0)  $\rightarrow$ PhaseDiagramResult

# disable\_global\_minimization()

Disables global minimization.

Default: Enabled

#### Returns

This PhaseDiagramCalculation object

# dont\_keep\_default\_equilibria()

Do not keep the initial equilibria added by default.

This is only relevant in combination with add\_initial\_equilibrium().

This is the default behavior.

# Returns

This PhaseDiagramCalculation object

# enable\_global\_minimization()

Enables global minimization.

# Default: Enabled

# Returns

This PhaseDiagramCalculation object

# get\_components() $\rightarrow$ List[str]

Returns the names of the components in the system (including all components auto-selected by the database(s)).

#### Returns

The component names

# get\_gibbs\_energy\_addition\_for(phase: str) $\rightarrow$ float

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

#### Parameters

phase – Specify the name of the (stoichiometric or solution) phase with the addition

#### Returns

Gibbs energy addition to G per mole formula unit.

# $\texttt{get\_system\_data()} \rightarrow \textit{SystemData}$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using *with\_system\_modifications()*.

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as \*.tdb-file.

#### Returns

The system data

## keep\_default\_equilibria()

Keep the initial equilibria added by default. This is only relevant in combination with *add\_initial\_equilibrium()*.

Default behavior is to not keep default equilibria.

# Returns

This PhaseDiagramCalculation object

#### remove\_all\_initial\_equilibria()

Removes all previously added initial equilibria.

# Returns

This PhaseDiagramCalculation object

# run\_poly\_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

# Parameters

command – The Thermo-Calc Console Mode command

#### Returns

This PhaseDiagramCalculation object

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

# set\_gibbs\_energy\_addition\_for(phase: str, gibbs\_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs\_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

#### Parameters

- phase Specify the name of the (stoichiometric or solution) phase with the addition
- gibbs\_energy Addition to G per mole formula unit

#### Returns

This PhaseDiagramCalculation object

#### set\_phase\_to\_dormant(phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

#### **Parameters**

phase - The phase name or ALL\_PHASES for all phases

#### Returns

This PhaseDiagramCalculation object

# set\_phase\_to\_entered(phase: str, amount: float = 1.0)

Sets the phase to the status ENTERED, that is the default state.

#### Parameters

- **phase** The phase name or *ALL\_PHASES* for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

#### Returns

This PhaseDiagramCalculation object

# set\_phase\_to\_fixed(phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

# **Parameters**

- **phase** The phase name
- **amount** The fixed phase fraction (between 0.0 and 1.0)

# Returns

This PhaseDiagramCalculation object

## set\_phase\_to\_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

#### Parameters

**phase** – The phase name or *ALL\_PHASES* for all phases

# Returns

This PhaseDiagramCalculation object

# with\_options(options: PhaseDiagramOptions)

Sets the simulation options.

**Parameters options** – The simulation options

Returns

This PhaseDiagramCalculation object

with\_reference\_state(component: str, phase: str = 'SER', temperature: float = -1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

# Parameters

- **component** The name of the element must be given.
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT\_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The Pressure (in Pa) for the reference state.

#### Returns

This PhaseDiagramCalculation object

#### with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

**Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

# **Parameters**

**system\_modifications** – The system modification to be performed

# Returns

This PhaseDiagramCalculation object

# class tc\_python.step\_or\_map\_diagrams.AbstractPropertyDiagramCalculation(calculator)

Bases: AbstractCalculation

Abstract configuration required for a property diagram calculation.

Note: This is an abstract class that cannot be used directly.

**abstract calculate**(*keep\_previous\_results: bool = False, timeout\_in\_minutes: float = 0.0*)

# disable\_global\_minimization()

Disables global minimization.

**Default**: Enabled

#### Returns

This PropertyDiagramCalculation object

# disable\_step\_separate\_phases()

Disables *step separate phases*. This is the **default** setting.

## Returns

This PropertyDiagramCalculation object

# enable\_global\_minimization()

Enables global minimization.

Default: Enabled

#### Returns

This PropertyDiagramCalculation object

# enable\_step\_separate\_phases()

Enables step separate phases.

Default: By default separate phase stepping is disabled

**Note:** This is an advanced option, it is used mostly to calculate how the Gibbs energy for a number of phases varies for different compositions. This is particularly useful to calculate Gibbs energies for complex phases with miscibility gaps and for an ordered phase that is never disordered (e.g. SIGMA-phase, G-phase, MU-phase, etc.).

# Returns

This PropertyDiagramCalculation object

# $\texttt{get\_components()} \rightarrow List[str]$

Returns the names of the components in the system (including all components auto-selected by the database(s)).

# Returns

The component names

# get\_gibbs\_energy\_addition\_for(phase: str) → float

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

# Parameters

phase – Specify the name of the (stoichiometric or solution) phase with the addition

#### Returns

Gibbs energy addition to G per mole formula unit.

#### get\_system\_data() $\rightarrow$ SystemData

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with\_system\_modifications().

Note: Parameters can only be read from unencrypted (i.e. *user*) databases loaded as \*.*tdb*-file.

#### Returns

The system data

#### run\_poly\_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

#### Parameters

command - The Thermo-Calc Console Mode command

#### Returns

This PropertyDiagramCalculation object

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

#### set\_gibbs\_energy\_addition\_for(phase: str, gibbs\_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs\_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

# Parameters

- phase Specify the name of the (stoichiometric or solution) phase with the addition
- gibbs\_energy Addition to G per mole formula unit

#### Returns

This PropertyDiagramCalculation object

#### set\_phase\_to\_dormant(phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

#### Parameters

phase - The phase name or ALL\_PHASES for all phases

#### Returns

This PropertyDiagramCalculation object

#### set\_phase\_to\_entered(phase: str, amount: float = 1.0)

Sets the phase to the status ENTERED, that is the default state.

#### Parameters

- phase The phase name or ALL\_PHASES for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

#### Returns

This PropertyDiagramCalculation object

# set\_phase\_to\_fixed(phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

# Parameters

- phase The phase name
- **amount** The fixed phase fraction (between 0.0 and 1.0)

#### Returns

This PropertyDiagramCalculation object

# set\_phase\_to\_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

# Parameters phase – The phase name or *ALL\_PHASES* for all phases

#### Returns

This PropertyDiagramCalculation object

# with\_options(options: PropertyDiagramOptions)

Sets the simulation options.

# Parameters

**options** – The simulation options

#### Returns

This PropertyDiagramCalculation object

# with\_reference\_state(component: str, phase: str = 'SER', temperature: float = -1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

#### **Parameters**

- component The name of the element must be given.
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT\_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The Pressure (in Pa) for the reference state.

#### Returns

This PropertyDiagramCalculation object

#### with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

**Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

#### Parameters

system\_modifications – The system modification to be performed

#### Returns

This PropertyDiagramCalculation object

# class tc\_python.step\_or\_map\_diagrams.AxisType

Bases: AbstractAxisType

Factory class providing objects for configuring a logarithmic or linear axis by using *AxisType.linear()* or *AxisType.logarithmic()*.

# classmethod linear()

Creates an object for configuring a linear calculation axis.

**Default**: A minimum number of 40 steps.

**Note:** The returned object can be configured regarding the maximum step size *or* the minimum number of steps on the axis.

Returns A new *Linear* object

#### classmethod logarithmic()

Creates an object for configuring a logarithmic calculation axis.

Default: A scale factor of 1.1

Note: The returned object can be configured regarding the scale factor.

#### Returns

A new Logarithmic object

class tc\_python.step\_or\_map\_diagrams.CalculationAxis(quantity: Union[ThermodynamicQuantity,

str])

Bases: object

A calculation axis used for property and phase diagram calculations.

**Note:** A calculation axis is defining the varied condition and the range of variation. It is the same concept as in Thermo-Calc *Graphical Mode* or *Console Mode*.

## Default: A Linear axis with a minimum number of 40 steps

set\_max(max: float)

Sets the maximum quantity value of the calculation axis.

# There is no default value set, it always needs to be defined.

#### **Parameters**

**max** – The maximum quantity value of the axis [unit according to the axis quantity]

#### Returns

This CalculationAxis object

# set\_min(min: float)

Sets the minimum quantity value of the calculation axis.

# There is no default value set, it always needs to be defined.

#### Parameters

min – The minimum quantity value of the axis [unit according to the axis quantity]

#### Returns

This CalculationAxis object

# set\_start\_at(at: float)

Sets the starting point of the calculation on the axis.

Default: The default starting point is the center between the minimum and maximum quantity value

#### **Parameters**

**at** – The starting point on the axis [unit according to the axis quantity]

# Returns

This CalculationAxis object

with\_axis\_type(axis\_type: AxisType)

Sets the type of the axis.

Default: A Linear axis with a minimum number of 40 steps

**Parameters** 

**axis\_type** – The axis type (linear or logarithmic)

# Returns

This CalculationAxis object

class tc\_python.step\_or\_map\_diagrams.Direction(value)

Bases: Enum

An enumeration.

 $DECREASE_FIRST_AXIS = 3$ 

 $DECREASE\_SECOND\_AXIS = 4$ 

**INCREASE\_FIRST\_AXIS = 0** 

**INCREASE\_SECOND\_AXIS = 1** 

# class tc\_python.step\_or\_map\_diagrams.InitialEquilibrium(first\_axis: float, second\_axis: float) Bases: object

# add\_equilibria\_at\_all\_phase\_changes()

This generates one start point for each set of phase change in the chosen direction of the specified axis This ensures finding all possible phase boundary lines (not just the first one) along such an axis direction.

Default behavior is to only generate one start point at the first phase change.

#### Returns

This InitialEquilibrium object

# add\_equilibria\_at\_first\_phase\_change()

This generates one start point at the first phase change.

This is the default behavior.

# Returns

This InitialEquilibrium object

#### set\_direction(direction enum: Direction)

Specifies along which axes the initial equilibria should be added.

The default direction is INCREASE FIRST AXIS.

# **Parameters** direction\_enum -

#### Returns

This InitialEquilibrium object

# class tc\_python.step\_or\_map\_diagrams.Linear

Bases: AxisType

Represents a linear axis.

# $get_type() \rightarrow str$

Convenience method for getting axis type.

Returns

The type

set\_max\_step\_size(max\_step\_size: float)

Sets the axis to use the *maximum step size* configuration.

Default: This is not the default which is minimum number of steps

Note: Either maximum step size or minimum number of steps can be used but not both at the same time.

#### **Parameters**

**max\_step\_size** – The maximum step size [unit according to the axis quantity]

# Returns

This Linear object

# set\_min\_nr\_of\_steps(min\_nr\_of\_steps: float = 40)

Sets the axis to use the *minimum number of steps* configuration.

Default: This is the default option (with a *minimum number of steps* of 40)

Note: Either maximum step size or minimum number of steps can be used but not both at the same time.

# Parameters min\_nr\_of\_steps – The minimum number of steps

#### Returns

This Linear object

**class** tc\_python.step\_or\_map\_diagrams.**Logarithmic**(*scale\_factor: float = 1.1*)

#### Bases: AxisType

Represents a logarithmic axis.

**Note:** A logarithmic axis is useful for low fractions like in a gas phase where 1E-7 to 1E-2 might be an interesting range. For the pressure a logarithmic axis is often also useful.

#### $get_type() \rightarrow str$

Convenience method for getting axis type.

Returns The type

# set\_scale\_factor(scale\_factor: float = 1.1)

Sets the scale factor.

Default: 1.1

## Parameters

**scale\_factor** – The scale factor setting the maximum factor between two calculated values, must be larger than 1.0

#### Returns

This Logarithmic object

#### class tc\_python.step\_or\_map\_diagrams.PhaseDiagramCalculation(calculator)

Bases: AbstractPhaseDiagramCalculation

Configuration for a phase diagram calculation.

Note: Specify the conditions, the calculation is performed with calculate().

#### add\_initial\_equilibrium(initial\_equilibrium: InitialEquilibrium)

Add initial equilibrium start points from which a phase diagram is calculated.

Scans along the axis variables and generates start points when the scan procedure crosses a phase boundary.

It may take a little longer to execute than using the minimum number of start points, as some lines may be calculated more than once. But the core remembers all node points and subsequently stops calculations along a line when it finds a known node point.

It is also possible to create a sequence of start points from one initial equilibria.

# Parameters initial\_equilibrium – The initial equilibrium

#### Returns

This PhaseDiagramCalculation object

 $\texttt{calculate}(\textit{keep\_previous\_results: bool = False, timeout\_in\_minutes: float = 0.0) \rightarrow \textit{PhaseDiagramResult}}$ 

Performs the phase diagram calculation.

**Warning:** If you use *keep\_previous\_results=True*, you must not use another calculator or even get results in between the calculations using *calculate()*. Then the previous results will actually be lost.

# Parameters

- **keep\_previous\_results** If True, results from any previous call to this method are appended. This can be used to combine calculations with multiple start points if the mapping fails at a certain condition.
- **timeout\_in\_minutes** Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

# Returns

A new *PhaseDiagramResult* object which later can be used to get specific values from the calculated result.

# disable\_global\_minimization()

Disables global minimization.

Default: Enabled

#### Returns

This PhaseDiagramCalculation object

# dont\_keep\_default\_equilibria()

Do not keep the initial equilibria added by default.

This is only relevant in combination with *add\_initial\_equilibrium()*.

This is the default behavior.

# Returns

This PhaseDiagramCalculation object

# enable\_global\_minimization()

Enables global minimization.

#### Default: Enabled

# Returns

This PhaseDiagramCalculation object

#### get\_components() $\rightarrow$ List[str]

Returns the names of the components in the system (including all components auto-selected by the database(s)).

# Returns

The component names

# $\texttt{get\_gibbs\_energy\_addition\_for}(\textit{phase: str}) \rightarrow \texttt{float}$

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

# Parameters

phase – Specify the name of the (stoichiometric or solution) phase with the addition

#### Returns

Gibbs energy addition to G per mole formula unit.

# $get_system_data() \rightarrow SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with\_system\_modifications().

**Note:** Parameters can only be read from unencrypted (i.e. *user*) databases loaded as \*.*tdb*-file.

#### Returns

The system data

# keep\_default\_equilibria()

Keep the initial equilibria added by default. This is only relevant in combination with *add\_initial\_equilibrium()*.

Default behavior is to not keep default equilibria.

# Returns

This PhaseDiagramCalculation object

# remove\_all\_conditions()

Removes all set conditions.

#### Returns

This PhaseDiagramCalculation object

# remove\_all\_initial\_equilibria()

Removes all previously added initial equilibria.

#### Returns

This PhaseDiagramCalculation object

remove\_condition(quantity: Union[ThermodynamicQuantity, str])

Removes the specified condition.

#### **Parameters**

**quantity** – The thermodynamic quantity to set as condition; a Console Mode syntax string can be used as an alternative (for example X(Cr))

Returns

This ThermodynamicCalculation object

#### run\_poly\_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

# Parameters

command – The Thermo-Calc Console Mode command

#### Returns

This PhaseDiagramCalculation object

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

#### set\_condition(quantity: Union[ThermodynamicQuantity, str], value: float)

Sets the specified condition.

#### **Parameters**

- **quantity** The thermodynamic quantity to set as condition; a Console Mode syntax string can be used as an alternative (for example *X*(*Cr*))
- value The value of the condition
- Returns

This PhaseDiagramCalculation object

# set\_gibbs\_energy\_addition\_for(phase: str, gibbs\_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs\_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

**Parameters** 

- phase Specify the name of the (stoichiometric or solution) phase with the addition
- gibbs\_energy Addition to G per mole formula unit

# Returns

This PhaseDiagramCalculation object

#### set\_phase\_to\_dormant(phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

#### Parameters

phase - The phase name or ALL\_PHASES for all phases

#### Returns

This PhaseDiagramCalculation object

#### set\_phase\_to\_entered(phase: str, amount: float = 1.0)

Sets the phase to the status ENTERED, that is the default state.

#### **Parameters**

- phase The phase name or ALL\_PHASES for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

#### Returns

This PhaseDiagramCalculation object

# set\_phase\_to\_fixed(phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

# Parameters

- phase The phase name
- **amount** The fixed phase fraction (between 0.0 and 1.0)

#### Returns

This PhaseDiagramCalculation object

# set\_phase\_to\_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

# **phase** – The phase name or *ALL\_PHASES* for all phases

**Parameters** 

Returns

This PhaseDiagramCalculation object

# with\_first\_axis(axis: CalculationAxis)

Sets the first calculation axis.

#### Parameters

**axis** – The axis

# Returns

This PhaseDiagramCalculation object

# with\_options(options: PhaseDiagramOptions)

Sets the simulation options.

# Parameters

**options** – The simulation options

Returns

This PhaseDiagramCalculation object

with\_reference\_state(component: str, phase: str = 'SER', temperature: float = -1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

#### **Parameters**

- component The name of the element must be given.
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT\_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The Pressure (in Pa) for the reference state.

#### Returns

This PhaseDiagramCalculation object

# with\_second\_axis(axis: CalculationAxis)

Sets the second calculation axis.

# Parameters

**axis** – The axis

# Returns

This PhaseDiagramCalculation object

# with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

**Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

#### **Parameters**

system\_modifications – The system modification to be performed

#### Returns

This PhaseDiagramCalculation object

#### class tc\_python.step\_or\_map\_diagrams.PhaseDiagramOptions

Bases: object

Simulation options for phase diagram calculations.

# disable\_approximate\_driving\_force\_for\_metastable\_phases()

Disables the approximation of the driving force for metastable phases.

Default: Enabled

**Note:** When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it is important that these driving forces are correct, use *disable\_approximate\_driving\_force\_for\_metastable\_pha* to force the calculation to converge for the metastable phases.

# Returns

This PhaseDiagramOptions object

#### disable\_control\_step\_size\_during\_minimization()

Disables stepsize control during minimization (non-global).

#### Default: Enabled

#### Returns

This PhaseDiagramOptions object

# disable\_force\_positive\_definite\_phase\_hessian()

Disables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

#### Default: Enabled

# Returns

This PhaseDiagramOptions object

# dont\_use\_auto\_start\_points()

Switches the usage of automatic starting points for the mapping off.

Default: Switched on

# Returns

This PhaseDiagramOptions object

# dont\_use\_inside\_mesh\_points()

Switches the usage of inside meshing points for the mapping off.

#### Default: Switched off

# Returns

This PhaseDiagramOptions object

# enable\_approximate\_driving\_force\_for\_metastable\_phases()

Enables the approximation of the driving force for metastable phases.

Default: Enabled

**Note:** When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it is important that these driving forces are correct, use *disable\_approximate\_driving\_force\_for\_metastable\_pha* to force the calculation to converge for the metastable phases.

# Returns

This PhaseDiagramOptions object

# enable\_control\_step\_size\_during\_minimization()

Enables stepsize control during normal minimization (non-global).

Default: Enabled

Returns

This PhaseDiagramOptions object

# enable\_force\_positive\_definite\_phase\_hessian()

Enables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

Default: Enabled

# Returns

This PhaseDiagramOptions object

#### set\_global\_minimization\_max\_grid\_points(max\_grid\_points: int = 2000)

Sets the maximum number of grid points in global minimization. \*\* Only applicable if global minimization is actually used\*\*.

Default: 2000 points

#### **Parameters**

max\_grid\_points - The maximum number of grid points

# Returns

This PhaseDiagramOptions object

#### set\_global\_minimization\_test\_interval(global\_test\_interval: int = 0)

Sets the interval for the global test.

# Default: 0

```
Parameters
global_test_interval – The global test interval
```

#### Returns

This PhaseDiagramOptions object

# set\_max\_no\_of\_iterations(max\_no\_of\_iterations: int = 500)

Set the maximum number of iterations.

Default: max. 500 iterations

**Note:** As some models give computation times of more than 1 CPU second/iteration, this number is also used to check the CPU time and the calculation stops if 500 CPU seconds/iterations are used.

Parameters
<b>max_no_of_iterations</b> – The max. number of iterations
Returns

This PhaseDiagramOptions object

set\_no\_of\_mesh\_along\_axis(no\_of\_mesh\_along\_axis: int = 3)

Sets the number of meshes along an axis for the mapping.

Default: 3

Parameters
 no\_of\_mesh\_along\_axis - The number of meshes

Returns

This PhaseDiagramOptions object

set\_required\_accuracy(accuracy: float = 1e-06)

Sets the required relative accuracy.

Default: 1.0E-6

**Note:** This is a relative accuracy, and the program requires that the relative difference in each variable must be lower than this value before it has converged. A larger value normally means fewer iterations but less accurate solutions. The value should be at least one order of magnitude larger than the machine precision.

#### **Parameters**

**accuracy** – The required relative accuracy

Returns

This PhaseDiagramOptions object

#### set\_smallest\_fraction(smallest\_fraction: float = 1e-12)

Sets the smallest fraction for constituents that are unstable.

It is normally only in the gas phase that you can find such low fractions.

The **default value** for the smallest site-fractions is 1E-12 for all phases except for IDEAL phase with one sublattice site (such as the GAS mixture phase in many databases) for which the default value is always as 1E-30.

# Parameters

smallest\_fraction – The smallest fraction for constituents that are unstable

#### Returns

This PhaseDiagramOptions object

# use\_auto\_start\_points()

Switches the usage of automatic starting points for the mapping on.

# Default: Switched on

#### Returns

This PhaseDiagramOptions object

# use\_inside\_mesh\_points()

Switches the usage of inside meshing points for the mapping off.

Default: Switched off

Returns

This PhaseDiagramOptions object

# class tc\_python.step\_or\_map\_diagrams.PhaseDiagramResult(result)

Bases: AbstractResult

Result of a phase diagram calculation, it can be evaluated using quantities or Console Mode syntax.

# add\_coordinate\_for\_phase\_label(x: float, y: float)

Sets a coordinate in the result plot for which the stable phases will be evaluated and provided in the result data object. This can be used to plot the phases of a region into the phase diagram or just to programmatically evaluate the phases in certain regions.

Warning: This method takes coordinates of the **plot** axes and not of the calculation axis.

# Parameters

- **x** The coordinate of the first **plot** axis ("x-axis") [unit of the **plot** axis]
- y The coordinate of the second plot axis ("y-axis") [unit of the plot axis]

# Returns

This PhaseDiagramResult object

# get\_values\_grouped\_by\_quantity\_of( $x_quantity$ : Union[ThermodynamicQuantity, str], y\_quantity: Union[ThermodynamicQuantity, str]) $\rightarrow$ PhaseDiagramResultValues

Returns x-y-line data grouped by the multiple datasets of the specified quantities (for example in dependency of components). The available quantities can be found in the documentation of the factory class ThermodynamicQuantity. Usually the result data represents the phase diagram.

**Note:** The different datasets will contain *NaN*-values between different subsections and are not sorted (because they are unsortable due to their nature).

**Note:** Its possible to use functions as axis variables, either by using *ThermodynamicQuantity.user\_defined\_function*, or by using an expression that contains '='.

**Example** get\_values\_grouped\_by\_quantity\_of('T', ThermodynamicQuantity.user\_defined\_function('HM.T'))

**Example** get\_values\_grouped\_by\_quantity\_of('T', 'CP=HM.T')

# Parameters

- **x\_quantity** The first quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example '*T*'), or even a function (for example '*f*=*T*\*1.01')
- **y\_quantity** The second quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example '*NV*'), or even a function (for example '*CP=HM.T*')

#### Returns

The phase diagram data

# get\_values\_grouped\_by\_stable\_phases\_of( $x_quantity$ : Union[ThermodynamicQuantity, str], y\_quantity: Union[ThermodynamicQuantity, str]) $\rightarrow$

# **PhaseDiagramResultValues**

Returns x-y-line data grouped by the sets of "stable phases" (for example "LIQUID" or "LIQUID + FCC\_A1"). The available quantities can be found in the documentation of the factory class ThermodynamicQuantity. Usually the result data represents the phase diagram.

**Note:** The different datasets will contain *NaN*-values between different subsections and are not sorted (because they are unsortable due to their nature).

**Note:** Its possible to use functions as axis variables, either by using *ThermodynamicQuantity.user\_defined\_function*, or by using an expression that contains '='.

**Example** get\_values\_grouped\_by\_quantity\_of('T', ThermodynamicQuantity.user\_defined\_function('HM.T'))

**Example** get\_values\_grouped\_by\_quantity\_of('T', 'CP=HM.T')

#### **Parameters**

- **x\_quantity** The first quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example 'T'), or even a function (for example 'f=T\*1.01')
- **y\_quantity** The second quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example '*NV*'), or even a function (for example '*CP=HM.T*')

# Returns

The phase diagram data

# remove\_phase\_labels()

Erases all added coordinates for phase labels.

# Returns

This PhaseDiagramResult object

# save\_to\_disk(path: str)

Saves the result to disc. Note that a result is a folder, containing potentially many files. The result can later be loaded with load\_result\_from\_disk()

# Parameters

**path** – the path to the folder you want the result to be saved in. It can be relative or absolute.

#### Returns

this PhaseDiagramResult object

## set\_phase\_name\_style(phase\_name\_style\_enum: PhaseNameStyle = PhaseNameStyle.NONE)

Sets the style of the phase name labels that will be used in the result data object (constitution description, ordering description,  $\dots$ ).

Default: PhaseNameStyle.NONE

#### **Parameters**

phase\_name\_style\_enum - The phase name style

#### Returns

This PhaseDiagramResult object

```
class tc_python.step_or_map_diagrams.PhaseDiagramResultValues(phase_diagram_values_java)
```

Bases: object

Represents the data of a phase diagram.

#### $get_invariants() \rightarrow ResultValueGroup$

Returns the x- and y-datasets of all invariants in the phase diagram.

Note: The datasets will normally contain different sections separated by NaN-values.

#### Returns

The invariants dataset object

#### **get\_lines()** → Dict[str, *ResultValueGroup*]

Returns the x- and y-datasets of all phase boundaries in the phase diagram.

Note: The datasets will normally contain different sections separated by NaN-values.

#### Returns

Containing the phase boundary datasets with the *quantities* or *stable phases* as keys (depending on the used method to get the values)

#### $get_phase_labels() \rightarrow List[PhaseLabel]$

Returns the phase labels added for certain coordinates using PhaseDiagramResult. add\_coordinate\_for\_phase\_label().

#### Returns

The list with the phase label data (that contains plot coordinates and stable phases)

#### get\_tie\_lines() → *ResultValueGroup*

Returns the x- and y-datasets of all tie-lines in the phase diagram.

**Note:** The datasets will normally contain different sections separated by *NaN*-values.

# Returns

The tie-line dataset object

#### class tc\_python.step\_or\_map\_diagrams.PhaseLabel(phase\_label\_java)

Bases: object

Represents a phase label at a plot coordinate, i.e. the stable phases that are present at that plot coordinate.

# $get_text() \rightarrow str$

Accessor for the phase label :return: the phase label

# $get_x() \rightarrow List[float]$

Accessor for the x-value :return: the x value

```
\texttt{get}\_y(\texttt{)} \to \texttt{List}[\texttt{float}]
```

Accessor for the y-value :return: the y value

# class tc\_python.step\_or\_map\_diagrams.PhaseNameStyle(value)

Bases: Enum

The style of the phase names used in the labels.

ALL = 1

Adding ordering and constitution description.

# CONSTITUTION\_DESCRIPTION = 3

Adding only constitution description.

```
NONE = 0
```

Only the phase names.

# **ORDERING\_DESCRIPTION = 4**

Adding only ordering description.

# class tc\_python.step\_or\_map\_diagrams.PropertyDiagramCalculation(calculator)

Bases: AbstractPropertyDiagramCalculation

# **calculate**(*keep\_previous\_results: bool* = *False*, *timeout\_in\_minutes: float* = 0.0) $\rightarrow$ *PropertyDiagramResult*

Performs the property diagram calculation.

**Warning:** If you use *keep\_previous\_results=True*, you must not use another calculator or even get results in between the calculations using *calculate()*. Then the previous results will actually be lost.

# Parameters

- **keep\_previous\_results** If *True*, results from any previous call to this method are appended. This can be used to combine calculations with multiple start points if the stepping fails at a certain condition.
- **timeout\_in\_minutes** Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

# Returns

A new *PropertyDiagramResult* object which later can be used to get specific values from the calculated result

# disable\_global\_minimization()

Disables global minimization.

Default: Enabled

# Returns

This PropertyDiagramCalculation object

#### disable\_step\_separate\_phases()

Disables step separate phases. This is the default setting.

#### Returns

This PropertyDiagramCalculation object

# enable\_global\_minimization()

Enables global minimization.

Default: Enabled

#### Returns

This PropertyDiagramCalculation object

# enable\_step\_separate\_phases()

Enables step separate phases.

Default: By default separate phase stepping is disabled

**Note:** This is an advanced option, it is used mostly to calculate how the Gibbs energy for a number of phases varies for different compositions. This is particularly useful to calculate Gibbs energies for complex phases with miscibility gaps and for an ordered phase that is never disordered (e.g. SIGMA-phase, G-phase, MU-phase, etc.).

#### Returns

This PropertyDiagramCalculation object

# $\texttt{get\_components()} \rightarrow List[str]$

Returns the names of the components in the system (including all components auto-selected by the database(s)).

# Returns

The component names

#### get\_gibbs\_energy\_addition\_for(*phase: str*) → float

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

# Parameters

phase - Specify the name of the (stoichiometric or solution) phase with the addition

#### Returns

Gibbs energy addition to G per mole formula unit.

# get\_system\_data() $\rightarrow$ SystemData

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with\_system\_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as \*.tdb-file.

#### Returns

The system data

# remove\_all\_conditions()

Removes all set conditions.

#### Returns

This PropertyDiagramCalculation object

#### remove\_condition(quantity: Union[ThermodynamicQuantity, str])

Removes the specified condition.

#### **Parameters**

**quantity** – The thermodynamic quantity to set as condition; a Console Mode syntax string can be used as an alternative (for example X(Cr))

#### Returns

This PropertyDiagramCalculation object

#### run\_poly\_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

#### Parameters

command – The Thermo-Calc Console Mode command

# Returns

This PropertyDiagramCalculation object

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

#### set\_condition(quantity: Union[ThermodynamicQuantity, str], value: float)

Sets the specified condition.

#### **Parameters**

- **quantity** The thermodynamic quantity to set as condition; a Console Mode syntax string can be used as an alternative (for example *X*(*Cr*))
- value The value of the condition

#### Returns

This PropertyDiagramCalculation object

# set\_gibbs\_energy\_addition\_for(phase: str, gibbs\_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs\_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

#### Parameters

- phase Specify the name of the (stoichiometric or solution) phase with the addition
- gibbs\_energy Addition to G per mole formula unit

#### Returns

This PropertyDiagramCalculation object

#### set\_phase\_to\_dormant(phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

#### Parameters

phase - The phase name or ALL\_PHASES for all phases

# Returns

This PropertyDiagramCalculation object

# set\_phase\_to\_entered(phase: str, amount: float = 1.0)

Sets the phase to the status ENTERED, that is the default state.

#### **Parameters**

- **phase** The phase name or *ALL\_PHASES* for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

# Returns

This PropertyDiagramCalculation object

#### set\_phase\_to\_fixed(phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

# Parameters

- phase The phase name
- **amount** The fixed phase fraction (between 0.0 and 1.0)

#### Returns

This PropertyDiagramCalculation object

#### set\_phase\_to\_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

#### Parameters

phase - The phase name or ALL\_PHASES for all phases

#### Returns

This PropertyDiagramCalculation object

# with\_axis(axis: CalculationAxis)

Sets the calculation axis.

# Parameters

**axis** – The axis

# Returns

This PropertyDiagramCalculation object

# with\_options(options: PropertyDiagramOptions)

Sets the simulation options.

# Parameters

options – The simulation options

#### Returns

This PropertyDiagramCalculation object

# with\_reference\_state(component: str, phase: str = 'SER', temperature: float = -1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state. All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

# **Parameters**

- **component** The name of the element must be given.
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT\_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The Pressure (in Pa) for the reference state.

# Returns

This PropertyDiagramCalculation object

# with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

**Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

# **Parameters**

system\_modifications - The system modification to be performed

# Returns

This PropertyDiagramCalculation object

# class tc\_python.step\_or\_map\_diagrams.PropertyDiagramOptions

Bases: object

Simulation options for the property diagram calculations.

# disable\_approximate\_driving\_force\_for\_metastable\_phases()

Disables the approximation of the driving force for metastable phases.

# Default: Enabled

**Note:** When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it is important that these driving forces are correct, use *disable\_approximate\_driving\_force\_for\_metastable\_pha* to force the calculation to converge for the metastable phases.

#### Returns

This PropertyDiagramOptions object

# disable\_control\_step\_size\_during\_minimization()

Disables stepsize control during minimization (non-global).

**Default**: Enabled

Returns

This PropertyDiagramOptions object

# disable\_force\_positive\_definite\_phase\_hessian()

Disables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

Default: Enabled

Returns

This PropertyDiagramOptions object

# enable\_approximate\_driving\_force\_for\_metastable\_phases()

Enables the approximation of the driving force for metastable phases.

Default: Enabled

**Note:** When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it is important that these driving forces are correct, use *disable\_approximate\_driving\_force\_for\_metastable\_pha* to force the calculation to converge for the metastable phases.

#### Returns

This PropertyDiagramOptions object

#### enable\_control\_step\_size\_during\_minimization()

Enables stepsize control during normal minimization (non-global).

# Default: Enabled

Returns

This PropertyDiagramOptions object

# enable\_force\_positive\_definite\_phase\_hessian()

Enables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

# Default: Enabled

Returns

This PropertyDiagramOptions object

# set\_global\_minimization\_max\_grid\_points(max\_grid\_points: int = 2000)

Sets the maximum number of grid points in global minimization. Only applicable if global minimization is actually used.

Default: 2000 points

#### Parameters

max\_grid\_points - The maximum number of grid points

Returns

This PropertyDiagramOptions object

# set\_global\_minimization\_test\_interval(global\_test\_interval: int = 0)

Sets the interval for the global test.

Default: 0

# Parameters global\_test\_interval – The global test interval

Returns

This *PropertyDiagramOptions* object

set\_max\_no\_of\_iterations(max\_no\_of\_iterations: int = 500)

Set the maximum number of iterations.

**Default**: max. 500 iterations

**Note:** As some models give computation times of more than 1 CPU second/iteration, this number is also used to check the CPU time and the calculation stops if 500 CPU seconds/iterations are used.

```
Parameters
max_no_of_iterations – The max. number of iterations
```

Returns

This PropertyDiagramOptions object

# set\_required\_accuracy(accuracy: float = 1e-06)

Sets the required relative accuracy.

Default: 1.0E-6

**Note:** This is a relative accuracy, and the program requires that the relative difference in each variable must be lower than this value before it has converged. A larger value normally means fewer iterations but less accurate solutions. The value should be at least one order of magnitude larger than the machine precision.

Parameters

**accuracy** – The required relative accuracy

Returns

This PropertyDiagramOptions object

# set\_smallest\_fraction(smallest\_fraction: float = 1e-12)

Sets the smallest fraction for constituents that are unstable.

It is normally only in the gas phase that you can find such low fractions.

The **default value** for the smallest site-fractions is 1E-12 for all phases except for IDEAL phase with one sublattice site (such as the GAS mixture phase in many databases) for which the default value is always as 1E-30.

# Parameters

smallest\_fraction – The smallest fraction for constituents that are unstable

Returns

This PropertyDiagramOptions object

class tc\_python.step\_or\_map\_diagrams.PropertyDiagramResult(result)

# Bases: AbstractResult

Result of a property diagram. This can be used to query for specific values.

get\_values\_grouped\_by\_quantity\_of( $x_quantity: Union[$ ThermodynamicQuantity, str],  $y_quantity: Union[$ ThermodynamicQuantity, str],  $sort_and_merge: bool = True) \rightarrow Dict[str, ResultValueGroup]$ 

Returns x-y-line data grouped by the multiple datasets of the specified quantities (typically the phases). The available quantities can be found in the documentation of the factory class ThermodynamicQuantity.

**Note:** The different datasets might contain *NaN*-values between different subsections and might not be sorted **even if the flag `sort\_and\_merge` has been set** (because they might be unsortable due to their nature).

**Note:** Its possible to use functions as axis variables, either by using *ThermodynamicQuantity.user\_defined\_function*, or by using an expression that contains '='.

**Example** get\_values\_grouped\_by\_quantity\_of('T', ThermodynamicQuantity.user\_defined\_function('HM.T'))

**Example** get\_values\_grouped\_by\_quantity\_of('T', 'CP=HM.T')

#### **Parameters**

- **x\_quantity** The first quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example 'T'), or even a function (for example 'f=T\*1.01')
- **y\_quantity** The second quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example '*NV*'), or even a function (for example '*CP=HM.T*')
- **sort\_and\_merge** If *True*, the data is sorted and merged into as few subsections as possible (divided by *NaN*)

# Returns

Containing the datasets with the quantities as their keys

# $get\_values\_grouped\_by\_stable\_phases\_of(x\_quantity: Union[ThermodynamicQuantity, str], y\_quantity: Union[ThermodynamicQuantity, str], sort\_and\_merge: bool = True) \rightarrow Dict[str, ResultValueGroup]$

Returns x-y-line data grouped by the sets of "stable phases" (for example "LIQUID" or "LIQUID + FCC\_A1"). The available quantities can be found in the documentation of the factory class ThermodynamicQuantity.

**Note:** The different datasets might contain *NaN*-values between different subsections and different lines of an ambiguous dataset. They might not be sorted **even if the flag `sort\_and\_merge` has been set** (because they might be unsortable due to their nature).

**Note:** Its possible to use functions as axis variables, either by using *ThermodynamicQuantity.user\_defined\_function*, or by using an expression that contains '='.

**Example** get\_values\_grouped\_by\_quantity\_of('T', ThermodynamicQuantity.user\_defined\_function('HM.T'))

**Example** get\_values\_grouped\_by\_quantity\_of('T', 'CP=HM.T')

# **Parameters**

- **x\_quantity** The first quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example '*T*'), or even a function (for example '*f*=*T*\**1.01*')
- **y\_quantity** The second quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example '*NV*'), or even a function (for example '*CP=HM.T*')
- **sort\_and\_merge** If *True*, the data will be sorted and merged into as few subsections as possible (divided by *NaN*)

#### Returns

Containing the datasets with the quantities as their keys

# **get\_values\_of**(x\_quantity: Union[ThermodynamicQuantity, str], y\_quantity: Union[ThermodynamicQuantity, str]) $\rightarrow$ [List[float], List[float]]

Returns sorted x-y-line data without any separation. Use *get\_values\_grouped\_by\_quantity\_of()* or *get\_values\_grouped\_by\_stable\_phases\_of()* instead if you need such a separation. The available quantities can be found in the documentation of the factory class ThermodynamicQuantity.

**Note:** This method will always return sorted data without any *NaN*-values. If it is unsortable that might give data that is hard to interpret. In such a case you need to choose the quantity in another way or use one of the other methods. One example of this is to use quantities with *All*-markers, for example *MassFractionOfAComponent("All")*.

**Note:** Its possible to use functions as axis variables, either by using *ThermodynamicQuantity.user\_defined\_function*, or by using an expression that contains '='.

**Example** get\_values\_grouped\_by\_quantity\_of('T', ThermodynamicQuantity.user\_defined\_function('HM.T'))

**Example** get\_values\_grouped\_by\_quantity\_of('T', 'CP=HM.T')

#### Parameters

• **x\_quantity** – The first Thermodynamic quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example 'T') or even a function (for example 'f=T\*1.01')

• **y\_quantity** – The second Thermodynamic quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example '*NV*'), or even a function (for example '*CP=HM.T*')

# Returns

A tuple containing the x- and y-data in lists

# save\_to\_disk(path: str)

Saves the result to disc. Note that a result is a folder, containing potentially many files. The result can later be loaded with load\_result\_from\_disk()

#### Parameters

**path** – the path to the folder you want the result to be saved in. It can be relative or absolute.

#### Returns

this PropertyDiagramResult object

set\_phase\_name\_style(phase\_name\_style\_enum: PhaseNameStyle = PhaseNameStyle.NONE)

Sets the style of the phase name labels that will be used in the result data object (constitution description, ordering description,  $\dots$ ).

Default: PhaseNameStyle.NONE

# Parameters phase\_name\_style\_enum - The phase name style

#### Returns

This PropertyDiagramResult object

# 6.1.6 Module "diffusion"

# class tc\_python.diffusion.AbstractBoundaryCondition

Bases: object

The abstract base class for all boundary conditions.

# class tc\_python.diffusion.AbstractCalculatedGrid

Bases: AbstractGrid

The abstract base class for calculated grids.

# class tc\_python.diffusion.AbstractElementProfile

Bases: object

The abstract base class for all initial composition profile types.

# class tc\_python.diffusion.AbstractGrid

Bases: object

The abstract base class for all grids.

# class tc\_python.diffusion.AbstractSolver

Bases: object

Abstract base class for the solvers (Classic, Homogenization and Automatic).

# class tc\_python.diffusion.ActivityFluxFunction

Bases: BoundaryCondition

#### $get_type() \rightarrow str$

The type of the boundary condition.

#### Returns

The type

**set\_flux\_function**(*element\_name: str, f: str = '0', g: str = '1', n: float = 1.0, to\_time: float = 1.7976931348623157e+308*)

The flux for the independent components must be given in the format:

 $J = f(T,P,TIME) * (ACTIVITY^N - g(T,P,TIME))$ 

where f and g may be functions of time (TIME), temperature (T), and pressure (P), and N is an integer.

f and g must be expressed in DICTRA Console Mode syntax.

# **Parameters**

- element\_name The name of the element
- **f** the function f in the formula above
- **g** the function g in the formula above
- **n** the constant N in the formula above
- to\_time The max-time for which the flux function is used.

# class tc\_python.diffusion.AutomaticGrid

Bases: CalculatedGrid

Represents an automatic grid.

# $\texttt{get\_type()} \rightarrow \textit{str}$

Type of the grid.

#### Returns

The type

# with\_coarse\_grid()

Configures this automatic grid as COARSE and returns the object to allow method chaining.

# Returns

This AutomaticGrid object (configured as COARSE).

#### **with\_custom\_grid**(*max\_number\_points: int = 50, max\_geometric\_factor: float = 1.2*)

Configures this automatic grid as CUSTOM and optionally sets max number points and max geometric factor.

#### Parameters

- **max\_number\_points** (Optional) Max number of points for the CUSTOM grid. Default is 50.
- **max\_geometric\_factor** (Optional) Max geometric factor for the CUSTOM grid. Default is 1.2.

#### Returns

This AutomaticGrid object (configured as CUSTOM).

# Raises

ValueError – If max\_number\_points is not greater than 1.
# with\_fine\_grid()

Configures this automatic grid as FINE and returns the object to allow method chaining.

## Returns

This AutomaticGrid object (configured as FINE).

## with\_medium\_grid()

Configures this automatic grid as MEDIUM and returns the object to allow method chaining.

#### Returns

This AutomaticGrid object (configured as MEDIUM).

# class tc\_python.diffusion.AutomaticGridType(value)

Bases: Enum

An enumeration representing the types of automatic grid.

Attributes: COARSE: Represents a coarse grid. MEDIUM: Represents a medium grid. FINE: Represents a fine grid. CUSTOM: Represents a CUSTOM grid.

COARSE = 'COARSE'

CUSTOM = 'CUSTOM'

FINE = 'FINE'

MEDIUM = 'MEDIUM'

## class tc\_python.diffusion.AutomaticSolver

Bases: Solver

Solver using the *homogenization model* if any region has more than one phase, otherwise using the *classic model*.

Note: This is the default solver and recommended for most applications.

## $get_type() \rightarrow str$

The type of the solver.

Returns

The type

# set\_flux\_balance\_equation\_accuracy(accuracy: float = 1e-16)

Only valid if the :class: ClassicSolver` is actually used (i.e. not more than one phase in each region).

Sets the required accuracy during the solution of the flux balance equations. Default: 1.0e-16

Parameters accuracy – The required accuracy

Returns

A new AutomaticSolver object

## set\_tieline\_search\_variable\_to\_activity()

Only valid if the :class: ClassicSolver` is actually used (i.e. not more than one phase in each region).

Configures the solver to use the *activity of a component* to find the correct tie-line at the phase interface. Either activity or chemical potential are applied to reduce the degrees of freedom at the local equilibrium. **Default**: This is the default setting

#### Returns

A new AutomaticSolver object

### set\_tieline\_search\_variable\_to\_potential()

## Only valid if the :class: ClassicSolver` is actually used (i.e. not more than one phase in each region).

Configures the solver to use the *chemical potential of a component* to find the correct tie-line at the phase interface. Either activity or chemical potential are applied to reduce the degrees of freedom at the local equilibrium. **Default**: To use the activity

#### Returns

A new AutomaticSolver object

# class tc\_python.diffusion.BoundaryCondition

Bases: AbstractBoundaryCondition

Contains factory methods for the the different boundary conditions available.

#### classmethod activity\_flux\_function()

Factory method that creates a **new** activity-flux-function boundary condition.

This type of boundary condition is used to take into account the finite rate of a surface reaction.

The flux for the independent components must be given in the format:

 $J = f(T,P,TIME) * (ACTIVITY^N - g(T,P,TIME))$ 

where f and g may be functions of time (TIME), temperature (T), and pressure (P), and N is an integer.

f and g must be expressed in DICTRA Console Mode syntax.

**Note:** The activities are those with user-defined reference states. The function mass transfer coefficient is the mass transfer coefficient, activity of the corresponding species in the gas is the activity of the corresponding species in the gas and N is a stoichiometric coefficient.

**Note:** For more details see L. Sproge and J. Ågren, "Experimental and theoretical studies of gas consumption in the gas carburizing process" J. Heat Treat. 6, 9–19 (1988).

#### Returns

A new ActivityFluxFunction object

## classmethod closed\_system()

Factory method that creates a **new** closed-system boundary condition.

## Returns

A new ClosedSystem object

#### classmethod fix\_flux\_value()

Factory method that creates a **new** fix-flux-value boundary condition.

This type of boundary condition makes it possible to enter functions that yield the flux times the molar volume for the independent components. May be a function of time, temperature and pressure: J(T,P,TIME).

#### Returns

A new FixFluxValue object

classmethod fixed\_compositions(unit\_enum: Unit = Unit.MASS\_PERCENT)

Factory method that creates a new fixed-composition boundary condition.

#### **Parameters**

unit\_enum – The composition unit

## Returns

A new FixedCompositions object

# classmethod mixed\_zero\_flux\_and\_activity()

Factory method that creates a new mixed zero-flux and activity boundary condition

## Returns

A new MixedZeroFluxAndActivity object

# class tc\_python.diffusion.CalculatedGrid

Bases: AbstractCalculatedGrid

Factory class for grids generated by a mathematical series (automatic, linear, geometric, ...). Use tc\_python. diffusion.PointByPointGrid instead if you want to use an existing grid from experimental data or a previous calculation.

**Note:** A region must contain a number of grid points. The composition is only known at these grid points and the software assumes that the composition varies linearly between them. The amount and composition of all the phases present at a single grid point in a certain region are those given by thermodynamic equilibrium keeping the over-all composition at the grid point fixed.

## classmethod automatic()

Factory method that creates a new automatic grid with MEDIUM grid type, as default.

Returns

A new AutomaticGrid object

```
classmethod double_geometric(no_of_points: int = 50, lower_geometrical_factor: float = 1.1,
upper_geometrical_factor: float = 0.9)
```

Factory method that creates a **new** double geometric grid.

**Note:** Double geometric grids have a high number of grid points in the middle or at both ends of a region. One geometrical factor for the lower (left) and upper (right) half of the region need to specified. In both cases a geometrical factor of larger than one yields a higher density of grid points at the lower end of the half and vice versa for a factor smaller than one.

## Parameters

- **no\_of\_points** The number of points
- lower\_geometrical\_factor The geometrical factor for the left half
- upper\_geometrical\_factor The geometrical factor for the right half

#### Returns

A new DoubleGeometricGrid object

**classmethod** geometric(*no\_of\_points: int* = 50, *geometrical\_factor: float* = 1.1)

Factory method that creates a **new** geometric grid.

**Note:** A grid that yields a varying density of grid points in the region. A geometrical factor larger than one yields a higher density of grid points at the lower end of the region and a factor smaller than one yields a higher density of grid points at the upper end of the region.

Parameters

- no\_of\_points The number of points
- geometrical\_factor The geometrical factor

## Returns

A new *GeometricGrid* object

# classmethod linear(no\_of\_points: int = 50)

Factory method that creates a new equally spaced grid.

Parameters no\_of\_points – The number of points

Returns

A new LinearGrid object

# class tc\_python.diffusion.ClassicSolver

Bases: Solver

Solver using the Classic model.

**Note:** This solver **never switches** to the homogenization model even if it fails to converge. Use the *tc\_python*. *diffusion*. *AutomaticSolver* if necessary instead.

#### $get_type() \rightarrow str$

Convenience method for getting the type of the solver.

#### Returns

The type of the solver

# set\_flux\_balance\_equation\_accuracy(accuracy: float = 1e-16)

Sets the required accuracy during the solution of the flux balance equations. Default: 1.0e-16

## Parameters

**accuracy** – The required accuracy

#### Returns

A new ClassicSolver object

## set\_tieline\_search\_variable\_to\_activity()

Configures the solver to use the *activity of a component* to find the correct tie-line at the phase interface. Either activity or chemical potential are applied to reduce the degrees of freedom at the local equilibrium. **Default**: This is the default setting

## set\_tieline\_search\_variable\_to\_potential()

Configures the solver to use the *chemical potential of a component* to find the correct tie-line at the phase interface. Either activity or chemical potential are applied to reduce the degrees of freedom at the local equilibrium. **Default**: To use the activity

#### Returns

A new ClassicSolver object

# class tc\_python.diffusion.ClosedSystem

Bases: BoundaryCondition

Represents a boundary for a closed system.

## $get_type() \rightarrow str$

Convenience method for getting the type of the boundary condition.

#### Returns

The type of the boundary condition

```
class tc_python.diffusion.CompositionProfile(unit_enum: Unit = Unit.MASS_PERCENT)
```

Bases: object

Contains initial concentration profiles for the elements.

add(element\_name: str, profile: ElementProfile)

Adds a concentration profile for the specified element.

#### **Parameters**

• **element\_name** – The name of the element

• **profile** – The initial concentration profile

#### Returns

A CompositionProfile object

# class tc\_python.diffusion.ConstantProfile(value: float)

Bases: ElementProfile

Represents a constant initial concentration profile.

## $get_type() \rightarrow str$

The type of the element profile.

## Returns

The type

#### class tc\_python.diffusion.ContinuedDiffusionCalculation(calculation)

Bases: AbstractCalculation

Configuration for a diffusion calculation that is a continuation of a previous isothermal or non-isothermal diffusion calculation. It contains a subset of the settings possible in the original calculation.

Use set\_simulation\_time() to set a simulation time that is higher than the original calculation.

## $calculate(timeout_in_minutes: float = 0.0) \rightarrow DiffusionCalculationResult$

Runs the diffusion calculation.

# **Parameters**

**timeout\_in\_minutes** – Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a Unre-coverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

## Returns

A *DiffusionCalculationResult* which later can be used to get specific values from the calculated result

# set\_simulation\_time(simulation\_time: float)

Sets the simulation time.

#### Parameters

**simulation\_time** – The simulation time [s]

#### Returns

This DiffusionIsoThermalCalculation object

# 

Defines the boundary condition on the left edge of the system.

Default: A closed-system boundary condition.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

**Note:** You can specify time-dependent boundary conditions by calling with\_left\_boundary\_condition() many times, with different values of the "to" parameter.

## Examples:

- with\_left\_boundary\_condition(BoundaryCondition.closed\_system(), to=100)
- with\_left\_boundary\_condition(BoundaryCondition.mixed\_zero\_flux\_and\_activity().set\_activity\_for\_element("C", surface\_activity), to=500)
- with\_left\_boundary\_condition(BoundaryCondition.closed\_system())

This example sets an closed-system-boundary-condition from start up to 100s and a activity-boundarycondition from 100s to 500s and finally a closed-system-boundary-condition from 500s to the end of simulation.

#### **Parameters**

- boundary\_condition The boundary condition
- **to** The upper time-limit for boundary\_condition.

#### Returns

This DiffusionIsoThermalCalculation object

with\_options(options: Options, to: float = 1.7976931348623157e+308)

Sets the general simulation conditions.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

# Parameters

- options The general simulation conditions
- to The upper time-limit for options.

#### Returns

This DiffusionIsoThermalCalculation object

## 

Defines the boundary condition on the right edge of the system.

Default: A closed-system boundary condition

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

**Note:** You can specify time-dependent boundary conditions by calling with\_right\_boundary\_condition() many times, with different values of the "to" parameter.

Examples:

- with\_right\_boundary\_condition(BoundaryCondition.closed\_system(), to=100)
- with\_right\_boundary\_condition(BoundaryCondition.mixed\_zero\_flux\_and\_activity().set\_activity\_for\_element("C", surface\_activity), to=500)
- with\_right\_boundary\_condition(BoundaryCondition.closed\_system())

This example sets an closed-system-boundary-condition from start up to 100s and a activity-boundarycondition from 100s to 500s and finally a closed-system-boundary-condition from 500s to the end of simulation.

#### **Parameters**

- **boundary\_condition** The boundary condition
- to The upper time-limit for boundary\_condition.

## Returns

This DiffusionIsoThermalCalculation object

with\_solver(solver: Solver, to: float = 1.7976931348623157e+308)

Sets the solver to use (Classic, Homogenization or Automatic). Default is Automatic.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

## **Parameters**

- **solver** The solver to use
- to The upper time-limit for solver.

#### Returns

This DiffusionIsoThermalCalculation object

with\_timestep\_control(timestep\_control: TimestepControl, to: float = 1.7976931348623157e+308)

Sets the timestep control options.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

## Parameters

- timestep\_control The new timestep control options
- to The upper time-limit for timestep\_control.

#### Returns

This DiffusionIsoThermalCalculation object

#### class tc\_python.diffusion.DiffusionCalculationResult(result)

#### Bases: AbstractResult

Result of a diffusion calculation. This can be used to query for specific values. For details of the axis variables, search the Thermo-Calc help.

get\_mass\_fraction\_at\_lower\_interface(region: str, component: str) → [List[float], List[float]]

Returns the mass fraction of the specified component at the lower boundary of the specified region, in dependency of time.

## Parameters

- region The name of the region
- **component** The name of the component

## Returns

A tuple of two lists of floats (time [s], mass fraction of the specified component)

# $\texttt{get_mass_fraction_at\_upper\_interface(\textit{region: str, component: str})} \rightarrow [\texttt{List[float]}, \texttt{List[float]}]$

Returns the mass fraction of the specified component at the upper boundary of the specified region, in dependency of time.

# **Parameters**

- **region** The name of the region
- **component** The name of the component

# Returns

A tuple of two lists of floats (time [s], mass fraction of the specified component)

 $\label{eq:get_mass_fraction_of_component_at_time(component: str, time: Union[SimulationTime, float]) \rightarrow \\ [List[float], List[float]]$ 

Returns the mass fraction of the specified component at the specified time.

**Note:** Use the enum *tc\_python.diffusion.SimulationTime* to choose the first or the last timepoint of the simulation. A timepoint close to the last one should never be specified manually because the actual end of the simulation can slightly deviate.

# Parameters

- **component** The name of the component
- **time** The time [s]

## Returns

A tuple of two lists of floats (distance [m], mass fraction of component at the specified time)

 $get\_mass\_fraction\_of\_phase\_at\_time(phase: str, time: Union[SimulationTime, float]) \rightarrow [List[float]] \rightarrow List[float]]$ 

Returns the mass fraction of the specified phase.

**Note:** Use the enum *tc\_python.diffusion.SimulationTime* to choose the first or the last timepoint of the simulation. A timepoint close to the last one should never be specified manually because the actual end of the simulation can slightly deviate.

# Parameters

- **phase** The name of the phase
- **time** The time [s]

# Returns

A tuple of two lists of floats (distance [m], mass fraction of hte phase at the specified time)

**get\_mole\_fraction\_at\_lower\_interface**(*region: str, component: str*)  $\rightarrow$  [List[float], List[float]]

Returns the mole fraction of the specified component at the lower boundary of the specified region, in dependency of time.

# Parameters

- region The name of the region
- **component** The name of the component

# Returns

A tuple of two lists of floats (time [s], mole fraction of the specified component)

# get\_mole\_fraction\_at\_upper\_interface(region: str, component: str) → [List[float], List[float]]

Returns the mole fraction of the specified component at the upper boundary of the specified region, in dependency of time.

## **Parameters**

- **region** The name of the region
- component The name of the component

## Returns

A tuple of two lists of floats (time [s], mole fraction of the specified component)

# $\texttt{get_mole_fraction_of_component_at_time}(\textit{component: str, time: Union[SimulationTime, float]}) \rightarrow [\texttt{List[float]},\texttt{List[float]}]$

Returns the mole fraction of the specified component at the specified time.

**Note:** Use the enum *tc\_python.diffusion.SimulationTime* to choose the first or the last timepoint of the simulation. A timepoint close to the last one should never be specified manually because the actual end of the simulation can slightly deviate.

# Parameters

- **component** The name of the component
- time The time [s]

## Returns

A tuple of two lists of floats (distance [m], mole fraction of component at the specified time)

 $\texttt{get\_mole\_fraction\_of\_phase\_at\_time(\textit{phase: str, time: Union[SimulationTime, float]}) \rightarrow [\texttt{List[float]}, \texttt{float}]) \rightarrow [\texttt{L$ 

```
List[float]]
```

Returns the mole fraction of the specified phase.

**Note:** Use the enum *tc\_python.diffusion.SimulationTime* to choose the first or the last timepoint of the simulation. A timepoint close to the last one should never be specified manually because the actual end of the simulation can slightly deviate.

# Parameters

• **phase** – The name of the phase

• **time** – The time [s]

#### Returns

A tuple of two lists of floats (distance [m], mole fraction of the phase at the specified time)

```
get_position_of_lower_boundary_of_region(region: str) → [List[float], List[float]]
```

Returns the position of the lower boundary of the specified region in dependency of time.

## Parameters

region – The name of the region

#### Returns

A tuple of two lists of floats (time [s], position of lower boundary of region [m])

# $\texttt{get_position_of\_upper\_boundary\_of\_region(\mathit{region: str}) \rightarrow [List[float], List[float]]$

Returns the position of the upper boundary of the specified region in dependency of time.

#### Parameters

**region** – The name of the region

## Returns

A tuple of two lists of floats (time [s], position of upper boundary of region [m])

# $\texttt{get\_regions()} \rightarrow List[str]$

Returns the regions of the diffusion simulation.

Note: Automatically generated regions  $(R_{\#})$  are included in the list.

#### Returns

The region names

# $\texttt{get\_time\_steps()} \rightarrow \texttt{List[float]}$

Returns the timesteps of the diffusion simulation.

## Returns

The timesteps [s]

# $\texttt{get\_total\_mass\_fraction\_of\_component}(\textit{component: str}) \rightarrow [\texttt{List[float]},\texttt{List[float]}]$

Returns the total mass fraction of the specified component in dependency of time.

# Parameters

component - The name of the component

## Returns

A tuple of two lists of floats (time [s], total mass fraction of the component)

get\_total\_mass\_fraction\_of\_component\_in\_phase(component: str, phase: str) → [List[float],

## List[float]]

Returns the total mass fraction of the specified component in the specified phase in dependency of time.

#### Parameters

- **component** The name of the component
- **phase** The name of the phase

#### Returns

A tuple of two lists of floats (time [s], total mass fraction of the component in the phase)

# get\_total\_mass\_fraction\_of\_phase(*phase: str*) → [List[float], List[float]]

Returns the total mass fraction of the specified phase in dependency of the time.

## Parameters

phase – The name of the phase

#### Returns

A tuple of two lists of floats (time [s], total mass fraction of the phase)

get\_total\_mole\_fraction\_of\_component(component: str) → [List[float], List[float]]

Returns the total mole fraction of the specified component in dependency of time.

## Parameters

component – The name of the component

#### Returns

A tuple of two lists of floats (time [s], total mole fraction of the component)

get\_total\_mole\_fraction\_of\_component\_in\_phase(component: str, phase: str)  $\rightarrow$  [List[float],

List[float]]

Returns the total mole fraction of the specified component in the specified phase in dependency of time.

## **Parameters**

- **component** The name of the component
- **phase** The name of the phase

#### Returns

A tuple of two lists of floats (time [s], total mole fraction of the component in the phase)

## get\_total\_mole\_fraction\_of\_phase(*phase: str*) → [List[float], List[float]]

Returns the total mole fraction of the specified phase in dependency of time.

#### Parameters

phase – The name of the phase

#### Returns

A tuple of two lists of floats (time [s], total mole fraction of the phase)

## get\_total\_volume\_fraction\_of\_phase(*phase: str*) → [List[float], List[float]]

Returns the total volume fraction of the specified phase in dependency of the time.

#### **Parameters**

**phase** – The name of the phase

#### Returns

A tuple of two lists of floats (time [s], total volume fraction of the phase)

 $get\_values\_of(x\_axis: Union[DiffusionQuantity, str], y\_axis: Union[DiffusionQuantity, str], plot\_condition: Union[PlotCondition, str] = ", independent\_variable: Union[IndependentVariable, str] = ") \rightarrow [List[float], List[float]]$ 

Returns the specified result from the simulation, allows all possible settings.

**Note:** As an alternative, DICTRA Console Mode syntax can be used as well for each quantity and condition.

**Warning:** This is an advanced mode that is equivalent to the possibilities in the DICTRA Console Mode. Not every combination of settings will return a result.

## Parameters

- **x\_axis** The first result quantity
- **y\_axis** The second result quantity
- plot\_condition The plot conditions
- independent\_variable The independent variable

#### Returns

A tuple of two lists of floats (the x\_axis quantity result, the y\_axis quantity result) [units according to the quantities]

## **get\_velocity\_of\_lower\_boundary\_of\_region**(*region: str*) $\rightarrow$ [List[float], List[float]]

Returns the velocity of the lower boundary of the specified region in dependency of time.

## **Parameters**

region – The name of the region

#### Returns

A tuple of two lists of floats (time [s], velocity of lower boundary of region [m/s])

# $\texttt{get\_velocity\_of\_upper\_boundary\_of\_region(\mathit{region: str}) \rightarrow [List[float], List[float]]$

Returns the velocity of the upper boundary of the specified region in dependency of time.

# **region** – The name of the region

#### Returns

**Parameters** 

A tuple of two lists of floats (time [s], velocity of upper boundary of region [m/s])

## **get\_width\_of\_region**(*region: str*) → [List[float], List[float]]

Returns the width of region, in dependency of time.

#### Parameters

 $\ensuremath{\textbf{region}}\xspace - \ensuremath{\textbf{The}}\xspace$  name of the region

#### Returns

A tuple of two lists of floats (time [s], width of the specified region [m])

## save\_to\_disk(path: str)

Saves the result to disk. The result can later be loaded using tc\_python.server.SetUp. load\_result\_from\_disk().

Note: The result data is represented by a whole folder containing multiple files.

## **Parameters**

**path** – The path to the result folder, can be relative or absolute.

# Returns

This DiffusionCalculationResult object

## with\_continued\_calculation()

Returns a *ContinuedDiffusionCalculation* that is used for continuing a diffusion calculation with altered settings.

#### Returns

A ContinuedDiffusionCalculation

## class tc\_python.diffusion.DiffusionIsoThermalCalculation(calculation)

Bases: AbstractCalculation

Configuration for an isothermal diffusion calculation.

# add\_console\_command(console\_command: str)

Registers a DICTRA Console Mode command for execution. These commands are executed after all other configuration directly before the calculation starts to run. All commands are stored and used until explicitly deleted using tc\_python.diffusion.DiffusionIsoThermoCalculation. remove\_all\_console\_commands.

#### **Parameters**

console\_command - The DICTRA Console Mode command

## Returns

This DiffusionIsoThermalCalculation object

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw DICTRA-commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten parenthesis, ...).

#### add\_region(region: Region)

Adds a region to the calculation. Regions are always added in the simulation domain from left to right.

If you want to replace an already added region, call *remove\_all\_regions()*, and add the regions that you want to keep.

Warning: Regions must have unique names.

Parameters

region-The region to be added

Returns

This DiffusionIsoThermalCalculation object

 $calculate(timeout_in_minutes: float = 0.0) \rightarrow DiffusionCalculationResult$ 

Runs the diffusion calculation.

#### **Parameters**

**timeout\_in\_minutes** – Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a Unre-coverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

#### Returns

A *DiffusionCalculationResult* which later can be used to get specific values from the calculated result

## $get_system_data() \rightarrow SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using *with\_system\_modifications()*.

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as \*.tdb-file.

Returns

The system data

## remove\_all\_console\_commands()

Removes all previously added Console Mode commands.

#### Returns

This DiffusionIsoThermalCalculation object

## remove\_all\_regions()

Removes all previously added regions.

:return This DiffusionIsoThermalCalculation object

# set\_simulation\_time(simulation\_time: float)

Sets the simulation time.

Parameters
 simulation\_time - The simulation time [s]

Returns

This DiffusionIsoThermalCalculation object

## set\_temperature(temperature: float)

Sets the temperature for the isothermal simulation.

Parameters temperature – The temperature [K]

Returns

This DiffusionIsoThermalCalculation object

## with\_cylindrical\_geometry(first\_interface\_position: float = 0.0)

Sets geometry to *cylindrical*, corresponds to an infinitely long cylinder of a certain radius.

Default: A planar geometry

**Note:** With a cylindrical or spherical geometry, the system's zero coordinate (left boundary) is at the centre of the cylinder or sphere by default. By specifying the *first\_interface\_position*, a different left-most coordinate can be defined. This allows to model a tube or a hollow sphere geometry. The highest coordinate (right boundary) is defined by the cylinder or sphere radius (i.e. by the width of all regions).

#### **Parameters**

**first\_interface\_position** – The position of the left-most coordinate along the axis, only necessary for modeling a tube geometry [m]

#### Returns

#### This DiffusionIsoThermalCalculation object

Defines the boundary condition on the left edge of the system.

Default: A closed-system boundary condition.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

**Note:** You can specify time-dependent boundary conditions by calling with\_left\_boundary\_condition() many times, with different values of the "to" parameter.

Examples:

- with\_left\_boundary\_condition(BoundaryCondition.closed\_system(), to=100)
- with\_left\_boundary\_condition(BoundaryCondition.mixed\_zero\_flux\_and\_activity().set\_activity\_for\_element("C", surface\_activity), to=500)
- with\_left\_boundary\_condition(BoundaryCondition.closed\_system())

This example sets an closed-system-boundary-condition from start up to 100s and a activity-boundarycondition from 100s to 500s and finally a closed-system-boundary-condition from 500s to the end of simulation.

## Parameters

- boundary\_condition The boundary condition
- **to** The upper time-limit for boundary\_condition.

#### Returns

This DiffusionIsoThermalCalculation object

with\_options(options: Options, to: float = 1.7976931348623157e+308)

Sets the general simulation conditions.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

#### Parameters

- options The general simulation conditions
- to The upper time-limit for options.

# Returns

This DiffusionIsoThermalCalculation object

#### with\_planar\_geometry()

Sets geometry to planar.

## This is default.

#### Returns

This DiffusionIsoThermalCalculation object

# with\_reference\_state(element: str, phase: str = 'SER', temperature: float = -1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

# Parameters

- **element** The name of the element
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT\_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The pressure (in Pa) for the reference state

## Returns

# This DiffusionIsoThermalCalculation object

# 

Defines the boundary condition on the right edge of the system.

**Default**: A closed-system boundary condition

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

**Note:** You can specify time-dependent boundary conditions by calling *with\_right\_boundary\_condition()* many times, with different values of the "to" parameter.

Examples:

- with\_right\_boundary\_condition(BoundaryCondition.closed\_system(), to=100)
- with\_right\_boundary\_condition(BoundaryCondition.mixed\_zero\_flux\_and\_activity().set\_activity\_for\_element("C", surface\_activity), to=500)
- with\_right\_boundary\_condition(BoundaryCondition.closed\_system())

This example sets an closed-system-boundary-condition from start up to 100s and a activity-boundarycondition from 100s to 500s and finally a closed-system-boundary-condition from 500s to the end of simulation.

## **Parameters**

- boundary\_condition The boundary condition
- to The upper time-limit for boundary\_condition.

#### Returns

This DiffusionIsoThermalCalculation object

with\_solver(solver: Solver, to: float = 1.7976931348623157e+308)

Sets the solver to use (Classic, Homogenization or Automatic). Default is Automatic.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

#### Parameters

- **solver** The solver to use
- to The upper time-limit for solver.

#### Returns

This DiffusionIsoThermalCalculation object

## with\_spherical\_geometry(first\_interface\_position: float = 0.0)

Sets geometry to *spherical*, corresponds to a sphere with a certain radius.

Default: A spherical geometry

**Note:** With a cylindrical or spherical geometry, the system's zero coordinate (left boundary) is at the centre of the cylinder or sphere by default. By specifying the *first\_interface\_position*, a different left-most coordinate can be defined. This allows to model a tube or a hollow sphere geometry. The highest coordinate (right boundary) is defined by the cylinder or sphere radius (i.e. by the width of all regions).

#### **Parameters**

**first\_interface\_position** – The position of the left-most coordinate along the axis, only necessary for modeling a hollow sphere geometry [m]

#### Returns

This DiffusionIsoThermalCalculation object

#### with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

**Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

Parameters

**system\_modifications** – The system modification to be performed

Returns

This DiffusionIsoThermalCalculation object

with\_timestep\_control(timestep\_control: TimestepControl, to: float = 1.7976931348623157e+308)
Sets the timestep control options.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

## Parameters

- **timestep\_control** The new timestep control options
- **to** The upper time-limit for timestep\_control.

### Returns

This DiffusionIsoThermalCalculation object

## class tc\_python.diffusion.DiffusionNonIsoThermalCalculation(calculation)

# Bases: AbstractCalculation

Configuration for a non-isothermal diffusion calculation.

## add\_console\_command(console\_command: str)

Registers a DICTRA Console Mode command for execution. These commands are executed after all other configuration directly before the calculation starts to run. All commands are stored and used until explicitly deleted using tc\_python.diffusion.DiffusionNonIsoThermalCalculation. remove\_all\_console\_commands.

#### **Parameters**

console\_command – The DICTRA Console Mode command

## Returns

This DiffusionNonIsoThermalCalculation object

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw DICTRA-commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten parenthesis, ...).

## add\_region(region: Region)

Adds a region to the calculation. Regions are always added in the simulation domain from left to right.

If you want to replace an already added region, call *remove\_all\_regions()*, and add the regions that you want to keep.

Warning: Regions must have unique names.

## Parameters

region - The region to be added

# Returns

This DiffusionNonIsoThermalCalculation object

# $calculate(timeout_in_minutes: float = 0.0) \rightarrow DiffusionCalculationResult$

Runs the diffusion calculation.

## Parameters

**timeout\_in\_minutes** – Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a Unre-coverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

## Returns

A *DiffusionCalculationResult* which later can be used to get specific values from the calculated result

# $get_system_data() \rightarrow SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using *with\_system\_modifications()*.

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as \*.tdb-file.

# Returns

The system data

# remove\_all\_console\_commands()

Removes all previously added Console Mode commands.

# Returns

This DiffusionNonIsoThermalCalculation object

# remove\_all\_regions()

Removes all previously added regions.

# Returns

This DiffusionNonIsoThermalCalculation object

## set\_simulation\_time(simulation\_time: float)

Sets the simulation time.

Parameters

simulation\_time - The simulation time [s]

## Returns

This DiffusionNonIsoThermalCalculation object

# with\_cylindrical\_geometry(first\_interface\_position: float = 0.0)

Sets geometry to cylindrical, corresponds to an infinitely long cylinder of a certain radius.

# Default: A planar geometry

**Note:** With a cylindrical or spherical geometry, the system's zero coordinate (left boundary) is at the centre of the cylinder or sphere by default. By specifying the *first\_interface\_position*, a different left-most coordinate can be defined. This allows to model a tube or a hollow sphere geometry. The highest coordinate (right boundary) is defined by the cylinder or sphere radius (i.e. by the width of all regions).

#### Parameters

**first\_interface\_position** – The position of the left-most coordinate along the axis, only necessary for modeling a tube geometry [m]

#### Returns

This DiffusionNonIsoThermalCalculation object

## 

Defines the boundary condition on the left edge of the system.

Default: A closed-system boundary condition.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

**Note:** You can specify time-dependent boundary conditions by calling *with\_left\_boundary\_condition()* many times, with different values of the "to" parameter.

## Examples:

- with\_left\_boundary\_condition(BoundaryCondition.closed\_system(), to=100)
- with\_left\_boundary\_condition(BoundaryCondition.mixed\_zero\_flux\_and\_activity().set\_activity\_for\_element("C", surface\_activity), to=500)
- with\_left\_boundary\_condition(BoundaryCondition.closed\_system())

This example sets an closed-system-boundary-condition from start up to 100s and a activity-boundarycondition from 100s to 500s and finally a closed-system-boundary-condition from 500s to the end of simulation.

## **Parameters**

- boundary\_condition The boundary condition
- **to** The upper time-limit for boundary\_condition.

#### Returns

This DiffusionNonIsoThermalCalculation object

with\_options(options: Options, to: float = 1.7976931348623157e+308)

Sets the general simulation conditions.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

#### **Parameters**

- **options** The general simulation conditions
- to The upper time-limit for options.

#### Returns

This DiffusionNonIsoThermalCalculation object

# with\_planar\_geometry()

Sets geometry to planar.

## This is default.

## Returns

#### This DiffusionNonIsoThermalCalculation object

# with\_reference\_state(element: str, phase: str = 'SER', temperature: float = -1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

#### **Parameters**

- **element** The name of the element
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT\_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The pressure (in Pa) for the reference state

#### Returns

This DiffusionNonIsoThermalCalculation object

## 

Defines the boundary condition on the right edge of the system.

**Default**: A closed-system boundary condition

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

**Note:** You can specify time-dependent boundary conditions by calling *with\_right\_boundary\_condition()* many times, with different values of the "to" parameter.

#### Examples:

- *with\_right\_boundary\_condition(BoundaryCondition.closed\_system(), to=100)*
- with\_right\_boundary\_condition(BoundaryCondition.mixed\_zero\_flux\_and\_activity().set\_activity\_for\_element("C", surface\_activity), to=500)

with\_right\_boundary\_condition(BoundaryCondition.closed\_system())

This example sets an closed-system-boundary-condition from start up to 100s and a activity-boundarycondition from 100s to 500s and finally a closed-system-boundary-condition from 500s to the end of simulation.

## **Parameters**

- **boundary\_condition** The boundary condition
- to The upper time-limit for boundary\_condition.

#### Returns

This DiffusionNonIsoThermalCalculation object

with\_solver(solver: Solver, to: float = 1.7976931348623157e+308)

Sets the solver to use (Classic, Homogenization or Automatic). Default is Automatic.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

# **Parameters**

- **solver** The solver to use
- **to** The upper time-limit for solver.

#### Returns

This DiffusionNonIsoThermalCalculation object

## with\_spherical\_geometry(first\_interface\_position: float = 0.0)

Sets geometry to *spherical*, corresponds to a sphere with a certain radius.

Default: A spherical geometry

**Note:** With a cylindrical or spherical geometry, the system's zero coordinate (left boundary) is at the centre of the cylinder or sphere by default. By specifying the *first\_interface\_position*, a different left-most coordinate can be defined. This allows to model a tube or a hollow sphere geometry. The highest coordinate (right boundary) is defined by the cylinder or sphere radius (i.e. by the width of all regions).

#### Parameters

**first\_interface\_position** – The position of the left-most coordinate along the axis, only necessary for modeling a hollow sphere geometry [m]

#### Returns

This DiffusionNonIsoThermalCalculation object

#### with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

**Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

**Parameters** 

system\_modifications - The system modification to be performed

### Returns

This DiffusionNonIsoThermalCalculation object

## with\_temperature\_profile(temperature\_profile: TemperatureProfile)

Sets the temperature profile to use with this calculation.

## Parameters

**temperature\_profile** – The temperature profile object (specifying time / temperature points)

#### Returns

This DiffusionNonIsoThermalCalculation object

with\_timestep\_control(timestep\_control: TimestepControl, to: float = 1.7976931348623157e+308)
Sets the timestep control options.

It is possible specify the upper time-point for which this setting is valid using the parameter "to".

Default: The end of the simulation.

#### **Parameters**

- timestep\_control The new timestep control options
- **to** The upper time-limit for timestep\_control.

## Returns

This DiffusionNonIsoThermalCalculation object

**class** tc\_python.diffusion.**DoubleGeometricGrid**(*no\_of\_points: int = 50, lower\_geometrical\_factor: float = 1.1, upper\_geometrical\_factor: float = 0.9*)

## Bases: CalculatedGrid

Represents a double geometric grid.

## get\_lower\_geometrical\_factor() $\rightarrow$ float

Returns the lower geometrical factor (for the left half).

## Returns

The lower geometrical factor

## get\_no\_of\_points() → int

Returns number of grid points.

#### Returns

The number of grid points

## $get_type() \rightarrow str$

Type of the grid.

# Returns

The type of the grid

## get\_upper\_geometrical\_factor()

Returns the upper geometrical factor (for the right half).

# Returns

The upper geometrical factor

## set\_lower\_geometrical\_factor(geometrical\_factor: float = 1.1)

Sets the lower (left half) geometrical factor.

**Note:** A geometrical factor of larger than one yields a higher density of grid points at the lower end of the half and vice versa for a factor smaller than one.

Parameters geometrical\_factor – The geometrical factor for the left half

Returns

This DoubleGeometricGrid object

## set\_no\_of\_points(no\_of\_points: int = 50)

Sets the number of grid points.

Parameters no\_of\_points – The number of points

Returns

This DoubleGeometricGrid object

## set\_upper\_geometrical\_factor(geometrical\_factor: float = 0.9)

Sets the upper (right half) geometrical factor.

**Note:** A geometrical factor of larger than one yields a higher density of grid points at the lower end of the half and vice versa for a factor smaller than one.

Parameters geometrical\_factor – The geometrical factor for the right half

#### Returns

This DoubleGeometricGrid object

## class tc\_python.diffusion.ElementProfile

Bases: AbstractElementProfile

Factory class providing objects for configuring a step, function or linear initial concentration profile.

# classmethod constant(value: float)

Factory method that creates a **new** constant initial concentration profile.

## Parameters

value – The constant composition in the region. [unit as defined in CompositionProfile].

#### Returns

A new ConstantProfile object

# classmethod funct(dictra\_console\_mode\_function: str)

Factory method that creates a **new** initial concentration profile defined by a function in DICTRA Console Mode syntax.

## **Parameters**

**dictra\_console\_mode\_function** – The function, expressed in DICTRA Console Mode syntax.

#### Returns

A new FunctionProfile object

**Note:** This is an advanced feature, preferably a complex concentration profile should be generated using third party libraries and added to the simulation using *tc\_python.diffusion.PointByPointGrid*.

#### classmethod linear(start\_value: float, end\_value: float)

Factory method that creates a **new** linear initial concentration profile.

## Parameters

- **start\_value** Composition at the left side of the region [unit as defined in *CompositionProfile*].
- **end\_value** Composition at the right side of the region [unit as defined in *CompositionProfile*].

#### Returns

A new LinearProfile object

## **classmethod step**(*lower\_boundary: float*, *upper\_boundary: float*, *step\_at: float*)

Factory method that creates a **new** initial concentration profile with a step at the specified distance, otherwise the composition is constant at the specified values.

#### Parameters

- **lower\_boundary** Composition before the step [unit as defined in *CompositionProfile*].
- upper\_boundary Composition after the step [unit as defined in CompositionProfile].
- step\_at The distance where the step should be [m].

# Returns

A new StepProfile object

# class tc\_python.diffusion.FixFluxValue

Bases: BoundaryCondition

# $\texttt{get\_type()} \to \mathsf{str}$

The type of the boundary condition.

Returns

The type

**set\_flux**(*element\_name: str*, *J: str* = '0', *to\_time: float* = 1.7976931348623157e+308)

Enter functions that yield the flux times the molar volume for the specified element. May be a function of time, temperature and pressure: J(T,P,TIME).

## Parameters

- element\_name The name of the element
- **J** the function *J*(*T*,*P*,*TIME*)
- to\_time The max-time for which the flux function is used.

## class tc\_python.diffusion.FixedCompositions(unit\_enum: Unit = Unit.MASS\_PERCENT)

# Bases: BoundaryCondition

Represents a boundary having fixed composition values.

## $get_type() \rightarrow str$

The type of the boundary condition.

# Returns

The type

# set\_composition(element\_name: str, value: float)

Sets the composition for the specified element.

Note: The boundary composition needs to be specified for each element.

# Parameters

- element\_name The name of the element
- value The composition value [unit according to the constructor parameter]

class tc\_python.diffusion.FunctionProfile(dictra\_console\_mode\_function: str)

Bases: ElementProfile

Creates an initial concentration profile defined by a function in DICTRA Console Mode syntax.

**Note:** This is an advanced feature, preferably a complex concentration profile should be generated using third party libraries and added to the simulation using *tc\_python.diffusion.PointByPointGrid*.

# get\_type() $\rightarrow$ str

The type of the element profile.

## Returns

The type

# class tc\_python.diffusion.GeneralLowerHashinShtrikman

Bases: HomogenizationFunctions

General lower Hashin-Shtrikman bounds: the outermost shell consists of the phase with the most sluggish kinetics.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

class tc\_python.diffusion.GeneralLowerHashinShtrikmanExcludedPhase(excluded\_phases: List[str] =

[])

## Bases: HomogenizationFunctions

General lower Hashin-Shtrikman bounds: the outermost shell consists of the phase with the most sluggish kinetics.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

The excluded phases are not considered when evaluating what phase has the most sluggish kinetics.

# class tc\_python.diffusion.GeneralUpperHashinShtrikman

Bases: HomogenizationFunctions

General upper Hashin-Shtrikman bounds: the innermost shell consists of the phase with the most sluggish kinetics.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

class tc\_python.diffusion.GeneralUpperHashinShtrikmanExcludedPhase(excluded\_phases: List[str] =
 [])

# Bases: HomogenizationFunctions

General upper Hashin-Shtrikman bounds: the innermost shell consists of the phase with the most sluggish kinetics.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

The excluded phases are not considered when evaluating what phase has the most sluggish kinetics.

#### **class** tc\_python.diffusion.**GeometricGrid**( $no_of_points$ : int = 50, geometrical\_factor: float = 1.1)

```
Bases: CalculatedGrid
```

Represents a geometric grid.

# $\texttt{get\_geometrical\_factor()} \rightarrow \texttt{float}$

Returns the geometrical factor.

#### Returns

The geometrical factor

# $get_no_of_points() \rightarrow int$

Returns the number of grid points.

#### Returns

The number of grid points

 $get_type() \rightarrow str$ 

Returns the type of grid.

## Returns The type

## set\_geometrical\_factor(geometrical\_factor: float = 1.1)

Sets the geometrical factor.

**Note:** A geometrical factor larger than one yields a higher density of grid points at the lower end of the region and a factor smaller than one yields a higher density of grid points at the upper end of the region.

**Parameters** geometrical\_factor – The geometrical factor

# Returns This GeometricGrid object

## set\_no\_of\_points(no\_of\_points: int = 50)

Sets the number of grid points.

# Parameters

**no\_of\_points** – The number of points

# Returns

This GeometricGrid object

# class tc\_python.diffusion.GridPoint(distance: float)

Bases: object

Represents a grid point, this is used in combination with grids of the type tc\_python.diffusion. PointByPointGrid.

# add\_composition(element: str, value: float)

Adds a composition for the specified element to the grid point.

# Parameters

• **element** – The element

• value – The composition value [unit as defined for the grid]

# Returns

This GridPoint object

# class tc\_python.diffusion.HashinShtrikmanBoundMajority

Bases: HomogenizationFunctions

Hashin-Shtrikman bounds with majority phase as matrix phase: the outermost shell consists of the phase with the highest local volume fraction.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

class tc\_python.diffusion.HashinShtrikmanBoundMajorityExcludedPhase(excluded\_phases: List[str]

=[])

# Bases: HomogenizationFunctions

Hashin-Shtrikman bounds with majority phase as matrix phase: the outermost shell consists of the phase with the highest local volume fraction.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

The excluded phases are not considered when evaluating what phase has the most sluggish kinetics.

# class tc\_python.diffusion.HashinShtrikmanBoundPrescribed(matrix\_phase: str)

Bases: HomogenizationFunctions

Hashin-Shtrikman bounds with prescribed phase as matrix phase: the outermost shell consists of the prescribed phase.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

class tc\_python.diffusion.HashinShtrikmanBoundPrescribedExcludedPhase(matrix\_phase: str,

excluded\_phases: List[str] = [])

Bases: HomogenizationFunctions

# class tc\_python.diffusion.HomogenizationFunction(value)

Bases: Enum

*Homogenization function* used for the *homogenization solver*. Many homogenization functions are based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase. **Default**: *RULE\_OF\_MIXTURES* (i.e. upper Wiener bounds)

## GENERAL\_LOWER\_HASHIN\_SHTRIKMAN = 0

General lower Hashin-Shtrikman bounds: the outermost shell consists of the phase with the most sluggish kinetics.

#### GENERAL\_UPPER\_HASHIN\_SHTRIKMAN = 1

General upper Hashin-Shtrikman bounds: the innermost shell consists of the phase with the most sluggish kinetics.

## HASHIN\_SHTRIKMAN\_BOUND\_MAJORITY = 2

Hashin-Shtrikman bounds with majority phase as matrix phase: the outermost shell consists of the phase with the highest local volume fraction.

## **INVERSE\_RULE\_OF\_MIXTURES = 4**

Lower Wiener bounds: the geometrical interpretation are continuous layers of each phase orthogonal to the direction of diffusion

#### RULE\_OF\_MIXTURES = 3

Upper Wiener bounds: the geometrical interpretation are continuous layers of each phase parallel with the direction of diffusion

## class tc\_python.diffusion.HomogenizationFunctions

Bases: object

## classmethod general\_lower\_hashin\_shtrikman()

Factory method that creates a **new** homogenization function of the type *GeneralLowerHashinShtrikman*.

General lower Hashin-Shtrikman bounds: the outermost shell consists of the phase with the most sluggish kinetics.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

#### Returns

A new GeneralLowerHashinShtrikman object

## classmethod general\_lower\_hashin\_shtrikman\_excluded\_phase(excluded\_phases: List[str] = [])

Factory method that creates a **new** homogenization function of the type *GeneralLowerHashinShtrikmanExcludedPhase*.

General lower Hashin-Shtrikman bounds: the outermost shell consists of the phase with the most sluggish kinetics.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase. The excluded phases are not considered when evaluating what phase has the most sluggish kinetics.

#### Parameters

excluded\_phases – The excluded phases

Returns

A new GeneralLowerHashinShtrikmanExcludedPhase object

#### classmethod general\_upper\_hashin\_shtrikman()

Factory method that creates a **new** homogenization function of the type *GeneralUpperHashinShtrikman*.

General upper Hashin-Shtrikman bounds: the innermost shell consists of the phase with the most sluggish kinetics.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

## Returns

A new GeneralUpperHashinShtrikman object

```
classmethod general_upper_hashin_shtrikman_excluded_phase(excluded_phases: List[str] = [])
```

Factory method that creates a **new** homogenization function of the type *GeneralUpperHashinShtrikmanExcludedPhase*.

General upper Hashin-Shtrikman bounds: the innermost shell consists of the phase with the most sluggish kinetics.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase. The excluded phases are not considered when evaluating what phase has the most sluggish kinetics.

#### Parameters

excluded\_phases – The excluded phases

## Returns

A new GeneralUpperHashinShtrikmanExcludedPhase object

## classmethod hashin\_shtrikman\_bound\_majority()

Factory method that creates a **new** homogenization function of the type *HashinShtrikmanBoundMajority*.

Hashin-Shtrikman bounds with majority phase as matrix phase: the outermost shell consists of the phase with the highest local volume fraction.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase.

#### Returns

A new HashinShtrikmanBoundMajority object

classmethod hashin\_shtrikman\_bound\_majority\_excluded\_phase(excluded\_phases: List[str] = [])

Factory method that creates a **new** homogenization function of the type *HashinShtrikmanBoundMajorityExcludedPhase*.

Hashin-Shtrikman bounds with majority phase as matrix phase: the outermost shell consists of the phase with the highest local volume fraction. Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase. The excluded phases are not considered when evaluating what phase has the most sluggish kinetics.

## Parameters

excluded\_phases - The excluded phases

Returns

A new HashinShtrikmanBoundMajorityExcludedPhase object

#### classmethod hashin\_shtrikman\_bound\_prescribed(matrix\_phase: str)

Factory method that creates a **new** homogenization function of the type *HashinShtrikmanBoundPrescribed*.

Hashin-Shtrikman bounds with prescribed phase as matrix phase: the outermost shell consists of the prescribed phase.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase. Parameters

**matrix\_phase** – The matrix phase

Returns

A new HashinShtrikmanBoundPrescribed object

classmethod hashin\_shtrikman\_bound\_prescribed\_excluded\_phase(matrix\_phase: str,

excluded\_phases: List[str] =

[])

Factory method that creates a **new** homogenization function of the type *HashinShtrikmanBoundPrescribedExcludedPhase*.

Hashin-Shtrikman bounds with prescribed phase as matrix phase: the outermost shell consists of the prescribed phase.

Based on a variant of the Hashin-Shtrikman bounds, their geometrical interpretation are concentric spherical shells of each phase. The excluded phases are not considered when evaluating what phase has the most sluggish kinetics.

Parameters

- matrix\_phase The matrix phase
- excluded\_phases The excluded phases

#### Returns

A new HashinShtrikmanBoundPrescribedExcludedPhase object

## classmethod inverse\_rule\_of\_mixtures()

Factory method that creates a **new** homogenization function of the type InverseRuleOfMixtures.

Lower Wiener bounds: the geometrical interpretation are continuous layers of each phase orthogonal to the direction of diffusion.

#### Returns

A new InverseRuleOfMixtures object

#### classmethod inverse\_rule\_of\_mixtures\_excluded\_phase(excluded\_phases: List[str] = [])

Factory method that creates a **new** homogenization function of the type *InverseRuleOfMixturesExcludedPhase*.

Lower Wiener bounds: the geometrical interpretation are continuous layers of each phase orthogonal to the direction of diffusion. Excluded phases are not considered in the diffusion calculations.

# Parameters

excluded\_phases – The excluded phases

Returns

A new InverseRuleOfMixturesExcludedPhase object

## classmethod labyrinth\_factor\_f(matrix\_phase: str)

Factory method that creates a **new** homogenization function of the type LabyrinthFactorF.

The labyrinth factor functions implies that all diffusion takes place in a single continuous matrix phase. The impeding effect on diffusion by phases dispersed in the matrix phase is taken into account by multiplying the flux with the volume fraction of the matrix phase.

#### Parameters

**matrix\_phase** – The matrix phase

## Returns

A new LabyrinthFactorF object

### classmethod labyrinth\_factor\_f2(matrix\_phase: str)

Factory method that creates a new homogenization function of the type LabyrinthFactorF2.

The labyrinth factor functions implies that all diffusion takes place in a single continuous matrix phase. The impeding effect on diffusion by phases dispersed in the matrix phase is taken into account by multiplying the flux with the volume fraction of the matrix phase squared.

#### Parameters

matrix\_phase - The matrix phase

Returns

A new LabyrinthFactorF2 object

## classmethod rule\_of\_mixtures()

Factory method that creates a **new** homogenization function of the type *RuleOfMixtures*.

Upper Wiener bounds: the geometrical interpretation are continuous layers of each phase parallel with the direction of diffusion.

#### Returns

A new RuleOfMixtures object

## classmethod rule\_of\_mixtures\_excluded\_phase(excluded\_phases: List[str] = [])

Factory method that creates a **new** homogenization function of the type *RuleOfMixturesExcludedPhase*.

Upper Wiener bounds: the geometrical interpretation are continuous layers of each phase parallel with the direction of diffusion. Excluded phases are not considered in the diffusion calculations.

```
Parameters
excluded_phases – The excluded phases
```

## Returns

A new RuleOfMixturesExcludedPhase object

#### class tc\_python.diffusion.HomogenizationSolver

Bases: Solver

Solver using the Homogenization model.

**Note:** This solver always uses the homogenization model, even if all regions have only one phase. The solver is **significantly slower than the Classic model**. Use the *tc\_python.diffusion.AutomaticSolver* instead if you do not need that behavior.

## disable\_global\_minimization()

Disables global minimization to be used in equilibrium calculations. Default: Disabled

**Note:** In general, using global minimization **significantly increases the simulation time**, but there is also a significantly reduced risk for non-converged equilibrium calculations.

#### Returns

A new HomogenizationSolver object

## disable\_interpolation\_scheme()

Configures the simulation not use *any interpolation scheme*. **Default**: To use the *logarithmic interpolation scheme* with 10000 discretization steps

**Note:** The homogenization scheme can be switched on by using *with\_linear\_interpolation\_scheme* or *with\_logarithmic\_interpolation\_scheme*.

## enable\_global\_minimization()

Enables global minimization to be used in equilibrium calculations. Default: Disabled

**Note:** In general, using global minimization **significantly increases the simulation time**, but there is also a significantly reduced risk for non-converged equilibrium calculations.

## Returns

A new HomogenizationSolver object

# $\texttt{get\_type()} \to \mathsf{str}$

The type of solver.

Returns

The type

## set\_fraction\_of\_free\_memory\_to\_use(fraction: float)

Sets the maximum fraction of free physical memory to be used by the interpolation scheme. **Default**: 1 / 10 of the free physical memory

## Parameters

fraction - The maximum free physical memory fraction to be used

#### Returns

A new HomogenizationSolver object

## set\_memory\_to\_use(memory\_in\_megabytes: float)

Sets the maximum physical memory in megabytes to be used by the interpolation scheme. **Default**: 1000 MBytes of the free physical memory

#### **Parameters**

memory\_in\_megabytes - The maximum physical memory to be used

#### Returns

A new HomogenizationSolver object

## with\_function(homogenization\_function: HomogenizationFunctions)

Sets the *homogenization function* used by the *homogenization model*.

## Parameters

 ${\bf homogenization\_function}$  – The homogenization function used by the homogenization model

#### Returns

A new HomogenizationSolver object

## with\_linear\_interpolation\_scheme(steps: int = 10000)

Configures the simulation to use the *linear interpolation scheme*. **Default**: To use the *logarithmic interpolation scheme* with 10000 discretization steps

#### Parameters

steps – The number of discretization steps in each dimension

## Returns

A new HomogenizationSolver object

## with\_logarithmic\_interpolation\_scheme(steps: int = 10000)

Configures the simulation to use the *linear interpolation scheme*. **Default**: To use the *logarithmic interpolation scheme* with 10000 discretization steps

#### Parameters

steps – The number of discretization steps in each dimension

Returns

A new HomogenizationSolver object

## class tc\_python.diffusion.InverseRuleOfMixtures

Bases: HomogenizationFunctions

Lower Wiener bounds: the geometrical interpretation are continuous layers of each phase orthogonal to the direction of diffusion.

#### class tc\_python.diffusion.InverseRuleOfMixturesExcludedPhase(excluded\_phases: List[str] = [])

## Bases: HomogenizationFunctions

Lower Wiener bounds: the geometrical interpretation are continuous layers of each phase orthogonal to the direction of diffusion.

Excluded phases are not considered in the diffusion calculations.

## class tc\_python.diffusion.LabyrinthFactorF(matrix\_phase: str)

Bases: HomogenizationFunctions

The labyrinth factor functions implies that all diffusion takes place in a single continuous matrix phase. The impeding effect on diffusion by phases dispersed in the matrix phase is taken into account by multiplying the flux with the volume fraction of the matrix phase.

## class tc\_python.diffusion.LabyrinthFactorF2(matrix\_phase: str)

Bases: HomogenizationFunctions

The labyrinth factor functions implies that all diffusion takes place in a single continuous matrix phase. The impeding effect on diffusion by phases dispersed in the matrix phase is taken into account by multiplying the flux with the volume fraction of the matrix phase squared.

## **class** tc\_python.diffusion.**LinearGrid**(*no\_of\_points: int = 50*)

Bases: CalculatedGrid

Represents an equally spaced grid.

#### get\_no\_of\_points() → int

Returns the number of grid points.

Returns

The number of grid points

get\_type()  $\rightarrow$  str

Type of the grid.

Returns The type set\_no\_of\_points(no\_of\_points: int = 50)
Sets the number of grid points.

Parameters

**no\_of\_points** – The number of points

Returns

This LinearGrid object

class tc\_python.diffusion.LinearProfile(start\_value: float, end\_value: float)

Bases: ElementProfile

Represents a linear initial concentration profile.

 $get_type() \rightarrow str$ 

The type of the element profile.

Returns

The type

class tc\_python.diffusion.MixedZeroFluxAndActivity

Bases: BoundaryCondition

Represents a boundary having zero-flux as well as fixed-activity conditions.

**Default**: On that boundary for every element without an explicitly defined condition, a zero-flux boundary condition is used.

 $get_type() \rightarrow str$ 

The type of the boundary condition.

Returns

The type

Sets an activity expression for an element at the boundary. Enter a formula that the software evaluates during the calculation.

The formula can be:

- a function of the variable TIME
- a constant

The formula must be written with these rules:

- a number must begin with a number (not a .)
- a number must have a dot or an exponent (*E*)

The operators +, -, \*, /, \*\* (exponentiation) can be used and with any level of parenthesis. As shown, the following operators must be followed by open and closed parentheses ()

- SQRT(X) is the square root
- *EXP*(*X*) is the exponential
- *LOG*(*X*) is the natural logarithm
- LOG10(X) is the base 10 logarithm
- SIN(X), COS(X), TAN(X), ASIN(X), ACOS(X), ATAN(X)
- SINH(X), COSH(X), TANH(X), ASINH(X), ACOSH(X), ATANH(X)

- SIGN(X)
- *ERF*(*X*) is the error function

Default: the expression entered is used for the entire simulation.

**Parameters** 

- element\_name The name of the element
- activity The activity
- to\_time The max-time for which the activity is used.

# set\_zero\_flux\_for\_element(element\_name: str)

Sets a zero-flux condition for an element at the boundary. **Default for all elements at the boundary** without an explicitly defined condition

#### Parameters

element\_name – The name of the element

## class tc\_python.diffusion.Options

Bases: object

General simulation conditions for the diffusion calculations.

# disable\_forced\_starting\_values\_in\_equilibrium\_calculations()

Disables forced starting values for the equilibrium calculations. The default is 'enable\_automatic\_forced\_starting\_values\_in\_equilibrium\_calculations'.

#### Returns

This Options object

# disable\_save\_results\_to\_file()

Disables the saving of results to file during the simulation. Default: Saving of the results at every timestep

#### Returns

This Options object

# enable\_automatic\_forced\_starting\_values\_in\_eq\_calculations()

Lets calculation engine decide if forced start values for the equilibrium calculations should be used. **This is the default setting**.

## Returns

This Options object

## enable\_forced\_starting\_values\_in\_equilibrium\_calculations()

Enables forced start values for the equilibrium calculations. The default is 'enable\_automatic\_forced\_starting\_values\_in\_equilibrium\_calculations'.

#### Returns

This Options object

## enable\_save\_results\_to\_file(every\_nth\_step: int = -1)

Enables and configures saving of results to file during the simulation. They can be saved for every n-th or optionally for every timestep (-1). **Default**: Saving of the results at every timestep

## Parameters

every\_nth\_step – -1 or a value ranging from 0 to 99

#### Returns

This Options object
## enable\_time\_integration\_method\_automatic()

Enables automatic selection of integration method. This is the default method.

#### Returns

This Options object

## enable\_time\_integration\_method\_euler\_backwards()

Enables *Euler backwards* integration. The default method is enable\_time\_integration\_method\_automatic.

**Note:** This method is more stable but less accurate and may be necessary if large fluctuations occur in the profiles.

#### Returns

This Options object

#### enable\_time\_integration\_method\_trapezoidal()

Enables trapezoidal integration.

**Note:** If large fluctuations occur in the profiles, it may be necessary to use the more stable but less accurate *Euler backwards method*.

#### Returns

This **Options** object

## 

Sets the default required driving force for phases allowed to form at the interfaces. Default: 1.0e-5

Note: The required driving force (evaluated as DGM(ph)) is used for determining whether an inactive phase is stable, i.e. actually formed. DGM represents the driving force normalized by RT and is dimensionless.

## Parameters

**driving\_force** – The driving force (*DGM*(*ph*)) [-]

## Returns

This *Options* object

## class tc\_python.diffusion.PointByPointGrid(unit\_enum: Unit = Unit.MASS\_PERCENT)

## Bases: AbstractGrid

Represents a point-by-point grid. This is setting the grid and the compositions at once, it is typically used to enter a measured composition profile or the result from a previous calculation.

Note: If a point-by-point grid is used, it is not necessary to specify the grid and composition profile separately.

## add\_point(grid\_point: GridPoint)

Adds a grid point to the grid.

Parameters grid\_point – The grid point

## Returns

This PointByPointGrid object

get\_type()  $\rightarrow$  str

Type of the grid.

Returns The type

## class tc\_python.diffusion.Region(name: str)

Bases: object

Represents a region of the simulation domain that can contain more that one phase.

**Note:** The first added phase represents the matrix phase, while all later added phases are *spheriod phases*, i.e. precipitate phases.

## add\_phase(phase\_name: str, is\_matrix\_phase: bool = False)

Adds a phase to the region, each region must contain at least one phase.

**Note:** Normally the *matrix phase* and the *precipitate phases* are automatically chosen based on the presence of all profile elements in the phase and if it has diffusion data. If multiple phases have equal properties, the phase that was added first is chosen. The matrix phase can be explicitly set by using *is\_matrix\_phase=True*.

**Note:** If multiple phases are added to a region, the *homogenization model* is applied. That means that average properties of the local phase mixture are used.

## **Parameters**

- phase\_name The phase name
- **is\_matrix\_phase** If set to *True* this phase is explicitly set as matrix phase for the region, if no phase is set to *True*, the matrix phase is chosen automatically

## Returns

This Region object

add\_phase\_allowed\_to\_form\_at\_left\_interface(phase\_name: str, driving\_force: float = 1e-05)

Adds a phase allowed to form at the left boundary of the region (an *inactive phase*). The phase will only appear at the interface as a new automatic region if the driving force to form it is sufficiently high.

## Parameters

- phase\_name The phase name
- **driving\_force** The driving force for the phase to form (*DGM*(*ph*))

#### Returns

This Region object

add\_phase\_allowed\_to\_form\_at\_right\_interface(phase\_name: str, driving\_force: float = 1e-05)

Adds a phase allowed to form at the right boundary of the region (an *inactive phase*). The phase will only appear at the interface as a new automatic region if the driving force to form it is sufficiently high.

## Parameters

• phase\_name – The phase name

• **driving\_force** – The driving force for the phase to form (*DGM*(*ph*))

## Returns

This Region object

## remove\_all\_phases()

Removes all previously added phases from the region.

## Returns

This Region object

## set\_width(width: float)

Defined the width of the region.

Note: This method needs only to be used if a calculated grid has been defined (using with\_grid()).

Parameters width – The width [m]

#### Returns

This Region object

## with\_composition\_profile(initial\_compositions: CompositionProfile)

Defines the initial composition profiles for all elements in the region.

**Note:** This method needs only to be used if a calculated grid has been defined (using *with\_grid()*).

## Parameters

initial\_compositions – The initial composition profiles for all elements

#### Returns

This Region object

## with\_grid(grid: CalculatedGrid)

Defines a calculated grid in the region. If measured composition profiles or the result from a previous calculation should be used, instead with\_point\_by\_point\_grid\_containing\_compositions() needs to be applied.

**Note:** The composition profiles need to be defined separately using *with\_composition\_profile()*, additionally the region width needs to be specified using *set\_width()*.

Parameters grid – The grid

#### Returns

This Region object

## with\_point\_by\_point\_grid\_containing\_compositions(grid: PointByPointGrid)

Defines a point-by-point grid in the region. This is setting the grid and the compositions at once, it is typically used to enter a measured composition profile or the result from a previous calculation. If the composition profile should be calculated (linear, geometric, ...) with grid() should be used instead.

**Note:** If a point-by-point grid is used, with\_grid(), with\_composition\_profile() and set\_width() are unnecessary and must not be used.

```
Parameters
```

**grid** – The point-by-point grid

## Returns

This Region object

#### class tc\_python.diffusion.RuleOfMixtures

Bases: HomogenizationFunctions

Upper Wiener bounds: the geometrical interpretation are continuous layers of each phase parallel with the direction of diffusion.

## **class** tc\_python.diffusion.**RuleOfMixturesExcludedPhase**(*excluded\_phases: List[str] = []*)

Bases: HomogenizationFunctions

Upper Wiener bounds: the geometrical interpretation are continuous layers of each phase parallel with the direction of diffusion.

Excluded phases are not considered in the diffusion calculations.

#### class tc\_python.diffusion.SimulationTime(value)

Bases: Enum

Specifying special time steps for the evaluation of diffusion results.

**Note:** These placeholders should be used because especially the actual last timestep will slightly differ from the specified end time of the simulation.

## FIRST = 0

Represents the first timestep of the simulation

## LAST = 1

Represents the last timestep of the simulation

#### class tc\_python.diffusion.Solver

Bases: AbstractSolver

Factory class providing objects representing a solver.

## classmethod automatic()

Factory method that creates a new *automatic solver*. This is the default solver and recommended for most applications.

**Note:** This solver uses the homogenization model if any region has more than one phase, otherwise it uses the classic model.

## Returns

A new AutomaticSolver object

## classmethod classic()

Factory method that creates a new classic solver.

**Note:** This solver never switches to the homogenization model even if the solver fails to converge. Use the *tc\_python.diffusion.AutomaticSolver* if necessary instead.

## Returns

A new ClassicSolver object

#### classmethod homogenization()

Factory method that creates a **new** homogenization solver.

**Note:** This solver always uses the homogenization model, even if all regions have only one phase. The solver is **significantly slower than the Classic model**. Use the  $tc_python.diffusion$ . *AutomaticSolver* instead if you do not need that behavior.

## Returns

A new HomogenizationSolver object

class tc\_python.diffusion.StepProfile(lower\_boundary: float, upper\_boundary: float, step\_at: float)

Bases: ElementProfile

Represents an initial constant concentration profile with a step at the specified position.

 $\texttt{get\_type()} \to \mathsf{str}$ 

The type of the element profile.

## Returns

The type

## class tc\_python.diffusion.TimestepControl

Bases: object

Settings that control the time steps in the simulation.

## disable\_check\_interface\_position()

Disables checking of the interface position, i.e. the timesteps are not controlled by the phase interface displacement during the simulation. The default setting is :func:`enable\_automatic\_check\_interface\_position`.

#### Returns

This TimestepControl object

#### enable\_automatic\_check\_interface\_position()

Lets calculation engine decide if checking of the interface position should be used. **This is the default setting**.

#### Returns

This TimestepControl object

## enable\_check\_interface\_position()

Enables checking of the interface position, i.e. the timesteps are controlled by the phase interface displacement during the simulation. **The default setting is :func:`enable\_automatic\_check\_interface\_position`**.

#### Returns

This TimestepControl object

```
set_initial_time_step(initial_time_step: float = 1e-07)
Sets the initial timestep. Default: 1.0e-7 s
```

Parameters
initial\_time\_step - The initial timestep [s]

#### Returns

This TimestepControl object

## set\_max\_absolute\_error(absolute\_error: float = 1e-05)

Sets the maximum absolute error. Default: 1.0e-5

## Parameters

**absolute\_error** – The maximum absolute error

#### Returns

This TimestepControl object

#### set\_max\_relative\_error(relative\_error: float = 0.05)

Sets the maximum relative error. Default: 0.05

## Parameters

relative\_error – The maximum relative error

#### Returns

This TimestepControl object

The maximum timestep allowed during the simulation, specified in percent of the simulation time. **Default**: 10.0%

#### Parameters

max\_timestep\_allowed\_as\_percent\_of\_simulation\_time - The maximum timestep
allowed [%]

#### Returns

This TimestepControl object

## set\_max\_timestep\_increase\_factor(max\_timestep\_increase\_factor: float = 2.0)

Sets the maximum timestep increase factor. Default: 2

Note: For example, if 2 is entered the maximum time step is twice as long as the previous time step taken.

#### Parameters

max\_timestep\_increase\_factor - The maximum timestep increase factor

#### Returns

This TimestepControl object

## set\_smallest\_time\_step\_allowed(smallest\_time\_step\_allowed: float = 1e-07)

Sets the smallest time step allowed during the simulation. This is required when using the automatic procedure to determine the time step. **Default**: 1.0e-7 s

#### **Parameters**

smallest\_time\_step\_allowed - The smalles timestep allowed [s]

Returns

This TimestepControl object

```
class tc_python.diffusion.Unit(value)
```

Bases: Enum

Represents a composition unit.

 $MASS_FRACTION = 2$ 

Mass fraction.

 $MASS_PERCENT = 3$ 

Mass percent.

 $MOLE_FRACTION = 0$ 

Mole fraction.

 $MOLE_PERCENT = 1$ 

Mole percent.

 $U_FRACTION = 4$ 

U fraction

## 6.1.7 Module "propertymodel"

## class tc\_python.propertymodel.PropertyModelCalculation(calculator)

Bases: AbstractCalculation

Configuration for a Property Model calculation.

Note: Specify the settings, the calculation is performed with calculate().

 $calculate(timeout_in_minutes: float = 0.0) \rightarrow PropertyModelResult$ 

Runs the Property Model calculation.

## **Parameters**

**timeout\_in\_minutes** – Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a CalculationEngineException will be thrown.

#### Returns

A PropertyModelResult which later can be used to get specific values from the simulation.

#### get\_argument\_default(*argument\_id: str*) → object

Returns the default value for the specified argument. The argument id can be obtained with get\_arguments().

#### Parameters

argument\_id - The argument id

#### Returns

The default value (the type depends on the argument)

## $\texttt{get\_argument\_description}(\textit{argument\_id: str}) \rightarrow \texttt{str}$

Returns the detailed description of the argument. The id can be obtained with get\_arguments().

Parameters argument\_id – The argument id

## Returns

The detailed description

## $get\_arguments() \rightarrow Set[str]$

Returns a list of the arguments of the Property Model.

**Note:** The arguments are the 'UI-panel components' defined in the Property Model interface method provide\_ui\_panel\_components(). They have the same id as specified in the Property Model. The naming is different because there is no UI present.

## Returns

The ids of the available arguments

## get\_dynamic\_arguments() $\rightarrow$ Set[str]

Returns a list of the dynamic arguments of the Property Model.

**Note:** Dynamic arguments are "extra" arguments created by pressing the "plus" button that can occur next to the UI-panel for some models, when running the Property Model from within Thermo-Calc. You can use them also from the API by *invoke\_dynamic\_argument()*.

#### Returns

The ids of the available dynamic arguments

## get\_model\_description() $\rightarrow$ str

Returns the description text of the current model.

Returns

the description

#### 

Returns the current value of an optimizable model parameter. The id can be obtained with *get\_model\_parameters()*.

#### Parameters

model\_parameter\_id - The model parameter id

#### Returns

The current value [unit according to the parameter meaning]

## get\_model\_parameters() $\rightarrow$ Set[str]

Returns a list of the optimizable model parameters.

**Note:** The model parameters are an optional set of variables that can be used within the Property Model. Typically they are used to provide the possibility to inject parameter values during an optimization into the model. This allows the dynamic development of Property Models that need to be fitted to experimental data. The model parameters are controlled with the Property Model interface methods *provide\_model\_parameters* and *set\_model\_parameter*.

#### Returns

The ids of the optimizable model parameters

#### $get_system_data() \rightarrow SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using *with\_system\_modifications()*.

Note: Parameters can only be read from unencrypted (i.e. *user*) databases loaded as \**.tdb*-file.

#### Returns

The system data

#### invoke\_dynamic\_argument(argument\_id: str)

Increases the number of instances of this dynamic argument by one, the argument will have an id such as *argument\_1*, *argument\_2*, ... if the dynamic argument is called *argument*.

Note: You can obtain all available dynamic arguments by using get\_dynamic\_arguments().

## **Parameters** argument\_id – argument\_id: The argument id

## Returns

This PropertyModelCalculation object

## remove\_all\_conditions()

Removes all set classic POLY conditions.

**Note:** This does not affect the compositions set by set\_composition().

## Returns

This PropertyModelCalculation object

## remove\_dependent\_element()

Removes a manually set dependent element. This method does not affect the automatic choice of the dependent element if *set\_composition()* is used.

## Returns

This PropertyModelCalculation object

#### set\_argument(argument: str, value: str)

Sets the specified model argument to the specified value. The id can be obtained with get\_arguments().

#### **Parameters**

- argument The argument id
- value The value [unit according to the argument meaning]

Returns

This PropertyModelCalculation object

#### set\_composition(element\_name: str, value: float)

Sets the composition of a element. The unit for the composition can be changed using set\_composition\_unit().

#### **Default**: Mole percent (CompositionUnit.MOLE\_PERCENT)

#### **Parameters**

- element\_name The element
- value The composition value [composition unit defined for the calculation]

#### Returns

This PropertyModelCalculation object

set\_composition\_unit(unit\_enum: CompositionUnit = CompositionUnit.MOLE\_PERCENT)

Sets the composition unit.

Default: Mole percent (CompositionUnit.MOLE\_PERCENT).

#### Parameters

unit\_enum – The new composition unit

#### Returns

This PropertyModelCalculation object

## set\_condition(classic\_condition: str, value: float)

Adds a classic POLY condition. If that method is used, all conditions need to be specified in such a way. If this method is used, it is necessary to set the dependent element manually using set\_dependent\_element().

Default if not specified: pressure P = 1e5 Pa, system size N = 1, Temperature T = 1000 K

Warning: It is not possible to mix POLY-commands and compositions using set\_composition().

**Note:** It should not be necessary for most users to use this method, try to use *set\_composition()* instead.

**Warning:** As this method runs raw POLY-commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten parenthesis, ...).

#### **Parameters**

• **classic\_condition** – The classic POLY condition (for example: *X*(*CR*))

• value – The value of the condition

## Returns

This PropertyModelCalculation object

## set\_dependent\_element(dependent\_element\_name: str)

Sets the dependent element manually.

**Note:** It should not be necessary for most users to use this method. Setting the dependent element manually is only necessary and allowed if *set\_condition()* is used.

#### **Parameters**

dependent\_element\_name - The name of the dependent element

#### Returns

This PropertyModelCalculation object

## set\_model\_parameter(model\_parameter\_id: str, value)

Resets an optimizable model parameter. The id can be obtained with get\_model\_parameters().

## Parameters

- model\_parameter\_id The model parameter id
- **value** The new value of the parameter

#### Returns

This PropertyModelCalculation object

## set\_temperature(temperature: float = 1000)

Sets the temperature.

**Default**: 1000 K

Parameters temperature – The temperature [K]

#### Returns

This PropertyModelCalculation object

## with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

**Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

**Parameters** 

system\_modifications - The system modification to be performed

## Returns

This PropertyModelCalculation object

#### class tc\_python.propertymodel.PropertyModelResult(result)

Bases: AbstractResult

The result of a Property Model calculation.

#### get\_result\_quantities() $\rightarrow$ Set[str]

Returns a list of the available result quantities defined in the Property Model.

#### Returns

The ids of the defined result quantities

## 

Returns the detailed description of the result quantity. The id can be obtained by *get\_result\_quantities()*.

#### Parameters

result\_quantity\_id - The result quantity id

#### Returns

The detailed description

## get\_single\_equilibrium\_result(result\_quantity\_id: str = ") $\rightarrow$ SingleEquilibriumResult

Returns a result quantity value. The available result quantities can be obtained by *get\_result\_quantities()*.

## Parameters

result\_quantity\_id – The id of the result quantity.

#### Returns

The requested value [unit depending on the quantity], if the result is a SingleEquilibriumResult, is returned.

## $get\_value\_of(\mathit{result\_quantity\_id: str}) \rightarrow Union[float, Dict[str, float]]$

Returns a result quantity value. The available result quantities can be obtained by get\_result\_quantities().

#### Parameters

result\_quantity\_id – The id of the result quantity

#### Returns

The requested value [unit depending on the quantity]. If the result is parameterized, parameter-value pairs are returned.

## save\_to\_disk(path: str)

Saves the result to disk. The result can later be loaded using tc\_python.server.SetUp. load\_result\_from\_disk().

Note: The *result data* is represented by a whole folder possibly containing multiple files.

#### Parameters

**path** – The path to the result folder, can be relative or absolute.

## Returns

This PropertyModelResult object

## 6.1.8 Module "material\_to\_material"

## class tc\_python.material\_to\_material.AbstractConstantCondition

Bases: object

The abstract base class for all constant conditions.

## class tc\_python.material\_to\_material.AbstractMaterialToMaterialCalculationAxis

Bases: object

The abstract base class of all calculation axis.

## class tc\_python.material\_to\_material.ConstantCondition

Bases: AbstractConstantCondition

A constant condition.

## classmethod fraction\_of\_material\_b(fraction\_of\_material\_b: float = 0.5)

Creates a constant fraction of material B condition object.

Note: The unit depends on the composition unit setting in the calculator object.

#### **Parameters**

fraction\_of\_material\_b - The fraction of material B [weight-fraction or mole-fraction]

#### Returns

The condition object

## classmethod temperature(temperature: float = 1000)

Creates a constant temperature condition object.

Parameters temperature – The temperature [K]

Returns

The condition object

Bases: MaterialToMaterialCalculationAxis

A fraction of material B axis.

class tc\_python.material\_to\_material.FractionOfMaterialBCondition(fraction\_of\_material\_b: float

= 0.5)

Bases: ConstantCondition

A constant fraction of material B condition.

## class tc\_python.material\_to\_material.MaterialToMaterialCalculationAxis

Bases: AbstractMaterialToMaterialCalculationAxis

A calculation axis.

**classmethod fraction\_of\_material\_b**(*from\_fraction: float = 0.0, to\_fraction: float = 1.0, start\_fraction: float = 0.5*)

Creates a fraction of material B axis object.

Note: The unit depends on the composition unit setting in the calculator.

#### **Parameters**

- from\_fraction The left axis limit [weight-fraction or mole-fraction]
- **to\_fraction** The right axis limit [weight-fraction or mole-fraction]
- start\_fraction The start fraction of the calculation [weight-fraction or mole-fraction]

#### Returns

A new FractionOfMaterialBAxis axis object

**classmethod temperature**(*from\_temperature: float = 1000, to\_temperature: float = 3000, start\_temperature: float = 2000*)

Creates a temperature calculation axis object.

**Parameters** 

- **from\_temperature** The left axis limit [K]
- to\_temperature The right axis limit [K]
- start\_temperature The start temperature of the calculation [K]

#### Returns

A new TemperatureAxis condition object

## class tc\_python.material\_to\_material.MaterialToMaterialCalculationContainer(instance)

Bases: object

Provides access to the calculation objects for all Material to Material calculations.

These are specialised calculations for mixtures of two materials A and B. Otherwise they behave identical to the corresponding regular single equilibrium, property diagram and phase diagram calculations.

with\_phase\_diagram\_calculation(default\_conditions: bool = True, components: List[str] = [])  $\rightarrow$ MaterialToMaterialPhaseDiagramCalculation

Creates a Material to Material phase diagram (map) calculation.

#### Parameters

- default\_conditions If *True*, automatically sets the conditions *N*=1 and *P*=100000
- **components** Specify here the components of the system (for example: [AL2O3, ...]), *only necessary if they differ from the elements*. If this option is used, **all elements** of the system need to be replaced by a component.

#### Returns

A new MaterialToMaterialPhaseDiagramCalculation object

with\_property\_diagram\_calculation(default\_conditions: bool = True, components: List[str] = [])  $\rightarrow$ MaterialToMaterialPropertyDiagramCalculation

Creates a Material to Material property diagram (step) calculation.

#### Parameters

- default\_conditions If *True*, automatically sets the conditions *N*=1 and *P*=100000
- **components** Specify here the components of the system (for example: [AL2O3, ...]), *only necessary if they differ from the elements*. If this option is used, **all elements** of the system need to be replaced by a component.

## Returns

A new MaterialToMaterialPropertyDiagramCalculation object

with\_single\_equilibrium\_calculation(default\_conditions: bool = True, components: List[str] = [])  $\rightarrow$  MaterialToMaterialSingleEquilibriumCalculation

Creates a Material to Material single equilibrium calculation.

#### **Parameters**

- default\_conditions If *True*, automatically sets the conditions *N*=1 and *P*=100000
- **components** Specify here the components of the system (for example: [AL2O3, ...]), *only necessary if they differ from the elements*. If this option is used, **all elements** of the system need to be replaced by a component.

#### Returns

A new MaterialToMaterialSingleEquilibriumCalculation object

# class tc\_python.material\_to\_material.MaterialToMaterialPhaseDiagramCalculation(calculator) Bases: AbstractPhaseDiagramCalculation

Configuration for a Material to Material phase diagram calculation.

Note: Specify the conditions, the calculation is performed with calculate().

## add\_initial\_equilibrium(initial\_equilibrium: InitialEquilibrium)

Add initial equilibrium start points from which a phase diagram is calculated.

Scans along the axis variables and generates start points when the scan procedure crosses a phase boundary.

It may take a little longer to execute than using the minimum number of start points, as some lines may be calculated more than once. But the core remembers all node points and subsequently stops calculations along a line when it finds a known node point.

It is also possible to create a sequence of start points from one initial equilibria.

## Parameters initial\_equilibrium – The initial equilibrium

Returns

This MaterialToMaterialPhaseDiagramCalculation object

**calculate**(keep\_previous\_results: bool = False, timeout\_in\_minutes: float = 0.0)  $\rightarrow$ MaterialToMaterialPhaseDiagramResult

Performs the phase diagram calculation.

**Warning:** If you use *keep\_previous\_results=True*, you must not use another calculator or even get results in between the calculations using *calculate()*. Then the previous results will actually be lost.

Parameters

- **keep\_previous\_results** If *True*, results from any previous call to this method are appended. This can be used to combine calculations with multiple start points if the mapping fails at a certain condition.
- **timeout\_in\_minutes** Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

#### Returns

A new *MaterialToMaterialPhaseDiagramResult* object which later can be used to get specific values from the calculated result.

## disable\_global\_minimization()

Disables global minimization.

Default: Enabled

#### Returns

This MaterialToMaterialPhaseDiagramCalculation object

## dont\_keep\_default\_equilibria()

Do not keep the initial equilibria added by default.

This is only relevant in combination with *add\_initial\_equilibrium()*.

This is the default behavior.

#### Returns

This MaterialToMaterialPhaseDiagramCalculation object

## enable\_global\_minimization()

Enables global minimization.

Default: Enabled

## Returns

This MaterialToMaterialPhaseDiagramCalculation object

#### get\_components() $\rightarrow$ List[str]

Returns the names of the components in the system (including all components auto-selected by the database(s)).

## Returns

The component names

#### get\_gibbs\_energy\_addition\_for(*phase: str*) → float

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

## **Parameters**

phase – Specify the name of the (stoichiometric or solution) phase with the addition

## Returns

Gibbs energy addition to G per mole formula unit.

## $get_system_data() \rightarrow SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using *with\_system\_modifications()*.

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as \*.tdb-file.

#### Returns

The system data

## keep\_default\_equilibria()

Keep the initial equilibria added by default. This is only relevant in combination with *add\_initial\_equilibrium()*.

Default behavior is to not keep default equilibria.

#### Returns

This MaterialToMaterialPhaseDiagramCalculation object

## remove\_all\_initial\_equilibria()

Removes all previously added initial equilibria.

## Returns

This MaterialToMaterialPhaseDiagramCalculation object

## run\_poly\_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

## Parameters

command – The Thermo-Calc Console Mode command

#### Returns

This MaterialToMaterialPhaseDiagramCalculation object

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

## set\_activities(activities: Dict[str, float])

Sets the constant activity conditions.

**Note:** The activity conditions are identical for both materials.

## Parameters

activities – The constant activities

## Returns

This MaterialToMaterialPhaseDiagramCalculation object

#### set\_composition\_unit(unit: CompositionUnit = CompositionUnit.MASS\_PERCENT)

Sets the composition unit of both materials A and B.

Default: Weight percent

#### Parameters

unit – The composition unit of both materials A and B

#### Returns

This MaterialToMaterialPhaseDiagramCalculation object

## set\_gibbs\_energy\_addition\_for(phase: str, gibbs\_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs\_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

## **Parameters**

• phase – Specify the name of the (stoichiometric or solution) phase with the addition

• gibbs\_energy – Addition to G per mole formula unit

#### Returns

This MaterialToMaterialPhaseDiagramCalculation object

**set\_material\_a**(composition: Dict[str, float], dependent\_component: Optional[str] = None)

Sets the composition of the material A.

The unit is set with set\_composition\_unit().

**Tip:** The material can also have constant activity conditions, they are set in *set\_activities()*.

#### **Parameters**

- composition The composition of the material A
- dependent\_component The dependent component of the material A

#### Returns

This MaterialToMaterialPhaseDiagramCalculation object

**set\_material\_b**(composition: Dict[str, float], dependent\_component: Optional[str] = None)

Sets the composition of the material B.

The unit is set with set\_composition\_unit().

Tip: The material can also have constant activity conditions, they are set in set\_activities().

#### **Parameters**

- composition The composition of the material B
- dependent\_component The dependent component of the material B

#### Returns

This MaterialToMaterialPhaseDiagramCalculation object

## set\_phase\_to\_dormant(phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

## Parameters

phase - The phase name or ALL\_PHASES for all phases

## Returns

This MaterialToMaterialPhaseDiagramCalculation object

## set\_phase\_to\_entered(phase: str, amount: float = 1.0)

Sets the phase to the status ENTERED, that is the default state.

## Parameters

- **phase** The phase name or *ALL\_PHASES* for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

## Returns

This MaterialToMaterialPhaseDiagramCalculation object

## set\_phase\_to\_fixed(phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

## Parameters

- **phase** The phase name
- **amount** The fixed phase fraction (between 0.0 and 1.0)

## Returns

This MaterialToMaterialPhaseDiagramCalculation object

## set\_phase\_to\_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

## Parameters

phase – The phase name or ALL\_PHASES for all phases

## Returns

This MaterialToMaterialPhaseDiagramCalculation object

## set\_pressure(pressure: float)

Sets the pressure (i.e. the condition P).

**Note:** If the flag *default\_conditions=True* has been set during the creation of the calculator, the pressure is set to 1000 hPa by default.

Parameters pressure – The pressure [Pa]

## Returns

This MaterialToMaterialPhaseDiagramCalculation object

#### set\_system\_size(system\_size: float)

Sets the system size (i.e. the condition 'N', the number of moles).

**Note:** If the flag *default\_conditions=True* has been set during the creation of the calculator, the system size is set to 1.0 moles by default.

#### **Parameters**

system\_size - The system size [mole]

#### Returns

This MaterialToMaterialPhaseDiagramCalculation object

## with\_first\_axis(axis: MaterialToMaterialCalculationAxis)

Sets the first axis (either temperature of fraction of material B). This calculation type requires that both temperature and fraction of material B axis are set.

#### **Parameters axis** – The axis

## Returns

This MaterialToMaterialPhaseDiagramCalculation object

## with\_options(options: PhaseDiagramOptions)

Sets the simulation options.

**Parameters options** – The simulation options

#### Returns

This PhaseDiagramCalculation object

## with\_reference\_state(component: str, phase: str = 'SER', temperature: float = -1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

#### **Parameters**

• **component** – The name of the element must be given.

- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT\_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The Pressure (in Pa) for the reference state.

#### Returns

This MaterialToMaterialPhaseDiagramCalculation object

#### with\_second\_axis(axis: MaterialToMaterialCalculationAxis)

Sets the second axis (either temperature of fraction of material B). This calculation type requires that both temperature and fraction of material B axis are set.

#### Parameters

**axis** – The axis

#### Returns

This MaterialToMaterialPhaseDiagramCalculation object

## with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

**Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

#### Parameters

system\_modifications – The system modification to be performed

#### Returns

This MaterialToMaterialPhaseDiagramCalculation object

## class tc\_python.material\_to\_material.MaterialToMaterialPhaseDiagramResult(result)

Bases: PhaseDiagramResult

Result of a Material to Material phase diagram calculation, it can be evaluated using quantities or Console Mode syntax.

## add\_coordinate\_for\_phase\_label(x: float, y: float)

Sets a coordinate in the result plot for which the stable phases will be evaluated and provided in the result data object. This can be used to plot the phases of a region into the phase diagram or just to programmatically evaluate the phases in certain regions.

Warning: This method takes coordinates of the plot axes and not of the calculation axis.

#### **Parameters**

- **x** The coordinate of the first **plot** axis ("x-axis") [unit of the **plot** axis]
- y The coordinate of the second plot axis ("y-axis") [unit of the plot axis]

#### Returns

This MaterialToMaterialPhaseDiagramResult object

## **get\_values\_grouped\_by\_quantity\_of**(x-quantity: Union[ThermodynamicQuantity, str], y\_quantity: Union[ThermodynamicQuantity, str]) $\rightarrow$ PhaseDiagramResultValues

Returns x-y-line data grouped by the multiple datasets of the specified quantities (for example in dependency of components). The available quantities can be found in the documentation of the factory class ThermodynamicQuantity. Usually the result data represents the phase diagram.

**Note:** The different datasets will contain *NaN*-values between different subsections and are not sorted (because they are unsortable due to their nature).

**Note:** Its possible to use functions as axis variables, either by using ThermodynamicQuantity. user\_defined\_function, or by using an expression that contains '='.

**Example** get\_values\_grouped\_by\_quantity\_of('T', ThermodynamicQuantity.user\_defined\_function('HM.T'))

**Example** get\_values\_grouped\_by\_quantity\_of('T', 'CP=HM.T')

## Parameters

- **x\_quantity** The first quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example 'T'), *MATERIAL\_B\_FRACTION*, or even a function (for example 'f=T\*1.01')
- **y\_quantity** The second quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example '*NV*'), *MATERIAL\_B\_FRACTION*, or even a function (for example '*CP=HM.T*')

#### Returns

The phase diagram data

## get\_values\_grouped\_by\_stable\_phases\_of( $x_quantity$ : Union[ThermodynamicQuantity, str], y\_quantity: Union[ThermodynamicQuantity, str]) $\rightarrow$ PhaseDiagramResultValues

Returns x-y-line data grouped by the sets of "stable phases" (for example "LIQUID" or "LIQUID + FCC\_A1"). The available quantities can be found in the documentation of the factory class ThermodynamicQuantity. Usually the result data represents the phase diagram.

**Note:** The different datasets will contain *NaN*-values between different subsections and are not sorted (because they are unsortable due to their nature).

**Note:** Its possible to use functions as axis variables, either by using ThermodynamicQuantity. user\_defined\_function, or by using an expression that contains '='.

Exampleget\_values\_grouped\_by\_quantity\_of('T',ThermodynamicQuantity\_user\_defined\_function('HM.T'))

**Example** get\_values\_grouped\_by\_quantity\_of('T', 'CP=HM.T')

Parameters

- **x\_quantity** The first quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example 'T'), *MATERIAL\_B\_FRACTION*, or even a function (for example 'f=T\*1.01')
- **y\_quantity** The second quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example '*NV*'), *MATERIAL\_B\_FRACTION*, or even a function (for example '*CP=HM.T*')

## Returns

The phase diagram data

## remove\_phase\_labels()

Erases all added coordinates for phase labels.

#### Returns

This MaterialToMaterialPhaseDiagramResult object

## save\_to\_disk(path: str)

Saves the result to disc. Note that a result is a folder, containing potentially many files. The result can later be loaded with load\_result\_from\_disk()

#### **Parameters**

**path** – the path to the folder you want the result to be saved in. It can be relative or absolute.

#### Returns

this MaterialToMaterialPhaseDiagramResult object

#### set\_phase\_name\_style(phase\_name\_style\_enum: PhaseNameStyle = PhaseNameStyle.NONE)

Sets the style of the phase name labels that will be used in the result data object (constitution description, ordering description,  $\dots$ ).

Default: PhaseNameStyle.NONE

Parameters phase\_name\_style\_enum – The phase name style

## Returns

This MaterialToMaterialPhaseDiagramResult object

## class tc\_python.material\_to\_material.MaterialToMaterialPropertyDiagramCalculation(calculator)

Bases: AbstractPropertyDiagramCalculation

Configuration for a Material to Material property diagram calculation.

**Note:** Specify the conditions and possibly other settings, the calculation is performed with *calculate()*.

**calculate**(keep\_previous\_results: bool = False, timeout\_in\_minutes: float = 0.0)  $\rightarrow$ MaterialToMaterialPropertyDiagramResult

Performs the Material to Material property diagram calculation.

**Warning:** If you use *keep\_previous\_results=True*, you must not use another calculator or even get results in between the calculations using *calculate()*. Then the previous results will actually be lost.

Parameters

- **keep\_previous\_results** If *True*, results from any previous call to this method are appended. This can be used to combine calculations with multiple start points if the stepping fails at a certain condition.
- **timeout\_in\_minutes** Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

#### Returns

A new *MaterialToMaterialPropertyDiagramResult* object which later can be used to get specific values from the calculated result

## disable\_global\_minimization()

Disables global minimization.

Default: Enabled

#### Returns

This MaterialToMaterialPropertyDiagramCalculation object

## disable\_step\_separate\_phases()

Disables *step separate phases*. This is the **default** setting.

#### Returns

This MaterialToMaterialPropertyDiagramCalculation object

## enable\_global\_minimization()

Enables global minimization.

Default: Enabled

#### Returns

This MaterialToMaterialPropertyDiagramCalculation object

## enable\_step\_separate\_phases()

Enables step separate phases.

Default: By default separate phase stepping is disabled

**Note:** This is an advanced option, it is used mostly to calculate how the Gibbs energy for a number of phases varies for different compositions. This is particularly useful to calculate Gibbs energies for complex phases with miscibility gaps and for an ordered phase that is never disordered (e.g. SIGMA-phase, G-phase, MU-phase, etc.).

#### Returns

This MaterialToMaterialPropertyDiagramCalculation object

## $\texttt{get\_components()} \rightarrow List[str]$

Returns the names of the components in the system (including all components auto-selected by the database(s)).

#### Returns

The component names

## get\_gibbs\_energy\_addition\_for(phase: str) → float

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

## Parameters

phase – Specify the name of the (stoichiometric or solution) phase with the addition

#### Returns

Gibbs energy addition to G per mole formula unit.

## $get_system_data() \rightarrow SystemData$

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using with\_system\_modifications().

Note: Parameters can only be read from unencrypted (i.e. *user*) databases loaded as \*.*tdb*-file.

#### Returns

The system data

#### run\_poly\_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

#### Parameters command – The Thermo-Calc Console Mode command

#### Returns

This MaterialToMaterialPropertyDiagramCalculation object

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

#### set\_activities(activities: Dict[str, float])

Sets the constant activity conditions.

Note: The activity conditions are identical for both materials.

Parameters

**activities** – The constant activities

## Returns

This MaterialToMaterialPropertyDiagramCalculation object

#### set\_composition\_unit(unit: CompositionUnit = CompositionUnit.MASS\_PERCENT)

Sets the composition unit of both materials A and B.

Default: Weight percent

#### Parameters

unit – The composition unit of both materials A and B

#### Returns

This MaterialToMaterialPropertyDiagramCalculation object

## set\_gibbs\_energy\_addition\_for(phase: str, gibbs\_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs\_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

## **Parameters**

• phase – Specify the name of the (stoichiometric or solution) phase with the addition

• gibbs\_energy – Addition to G per mole formula unit

#### Returns

This MaterialToMaterialPropertyDiagramCalculation object

set\_material\_a(composition: Dict[str, float], dependent\_component: Optional[str] = None)

Sets the composition of the material A.

The unit is set with set\_composition\_unit().

**Tip:** The material can also have constant activity conditions, they are set in *set\_activities()*.

#### **Parameters**

- composition The composition of the material A
- dependent\_component The dependent component of the material A

#### Returns

This MaterialToMaterialPropertyDiagramCalculation object

**set\_material\_b**(composition: Dict[str, float], dependent\_component: Optional[str] = None)

Sets the composition of the material B.

The unit is set with set\_composition\_unit().

Tip: The material can also have constant activity conditions, they are set in set\_activities().

#### **Parameters**

- composition The composition of the material B
- dependent\_component The dependent component of the material B

#### Returns

This MaterialToMaterialPropertyDiagramCalculation object

## set\_phase\_to\_dormant(phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

## Parameters

phase - The phase name or ALL\_PHASES for all phases

## Returns

This MaterialToMaterialPropertyDiagramCalculation object

## set\_phase\_to\_entered(phase: str, amount: float = 1.0)

Sets the phase to the status ENTERED, that is the default state.

## Parameters

- **phase** The phase name or *ALL\_PHASES* for all phases
- **amount** The phase fraction (between 0.0 and 1.0)

## Returns

This MaterialToMaterialPropertyDiagramCalculation object

## set\_phase\_to\_fixed(phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

## Parameters

- **phase** The phase name
- **amount** The fixed phase fraction (between 0.0 and 1.0)

## Returns

This MaterialToMaterialPropertyDiagramCalculation object

## set\_phase\_to\_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

## Parameters

phase – The phase name or ALL\_PHASES for all phases

## Returns

This MaterialToMaterialPropertyDiagramCalculation object

## set\_pressure(pressure: float)

Sets the pressure (i.e. the condition *P*).

**Note:** If the flag *default\_conditions=True* has been set during the creation of the calculator, the pressure is set to 1000 hPa by default.

Parameters pressure – The pressure [Pa]

## Returns

This MaterialToMaterialPropertyDiagramCalculation object

#### set\_system\_size(system\_size: float)

Sets the system size (i.e. the condition 'N', the number of moles).

**Note:** If the flag *default\_conditions=True* has been set during the creation of the calculator, the system size is set to 1.0 moles by default.

## Parameters

system\_size - The system size [mole]

#### Returns

This MaterialToMaterialPropertyDiagramCalculation object

#### with\_axis(axis: MaterialToMaterialCalculationAxis)

Sets the axis (either temperature of fraction of material B). This calculation type requires that either temperature or fraction of material B is set as a constant condition - the other one is set as an axis.

Parameters axis – The axis

## Returns

This MaterialToMaterialPropertyDiagramCalculation object

### with\_constant\_condition(condition: ConstantCondition)

Sets the constant condition (either temperature of fraction of material B). This calculation type requires that either temperature or fraction of material B is set as a constant condition - the other one is set as an axis.

Parameters condition – The condition

## Returns

This MaterialToMaterialPropertyDiagramCalculation object

## with\_options(options: PropertyDiagramOptions)

Sets the simulation options.

## Parameters

options – The simulation options

#### Returns

This MaterialToMaterialPropertyDiagramCalculation object

# with\_reference\_state(component: str, phase: str = 'SER', temperature: float = -1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

## Parameters

- component The name of the element must be given.
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT\_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- pressure The Pressure (in Pa) for the reference state.

#### Returns

This MaterialToMaterialPropertyDiagramCalculation object

## with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

**Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

#### **Parameters**

system\_modifications – The system modification to be performed

#### Returns

This MaterialToMaterialPropertyDiagramCalculation object

## class tc\_python.material\_to\_material.MaterialToMaterialPropertyDiagramResult(result)

## Bases: PropertyDiagramResult

Result of a Material to Material property diagram. It can be used to query for specific values.

get\_values\_grouped\_by\_quantity\_of( $x_quantity: Union[$ ThermodynamicQuantity, str],  $y_quantity: Union[$ ThermodynamicQuantity, str],  $sort_and_merge: bool = True) \rightarrow Dict[str, ResultValueGroup]$ 

Returns x-y-line data grouped by the multiple datasets of the specified quantities (typically the phases). The available quantities can be found in the documentation of the factory class ThermodynamicQuantity.

**Note:** The different datasets might contain *NaN*-values between different subsections and might not be sorted **even if the flag `sort\_and\_merge` has been set** (because they might be unsortable due to their nature).

**Note:** Its possible to use functions as axis variables, either by using ThermodynamicQuantity. user\_defined\_function, or by using an expression that contains '='.

**Example** get\_values\_grouped\_by\_quantity\_of('T', ThermodynamicQuantity.user\_defined\_function('HM.T')) **Example** get\_values\_grouped\_by\_quantity\_of('T', 'CP=HM.T')

## **Parameters**

- **x\_quantity** The first quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example '*T*'), *MATERIAL\_B\_FRACTION*, or even a function (for example '*f*=*T*\*1.01')
- **y\_quantity** The second quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example '*NV*'), *MATERIAL\_B\_FRACTION*, or even a function (for example '*CP=HM.T*')
- sort\_and\_merge If *True*, the data is sorted and merged into as few subsections as possible (divided by *NaN*)

#### Returns

Containing the datasets with the quantities as their keys

get\_values\_grouped\_by\_stable\_phases\_of(x-quantity: Union[ThermodynamicQuantity, str], y\_quantity: Union[ThermodynamicQuantity, str], sort\_and\_merge: bool = True)  $\rightarrow$  Dict[str, ResultValueGroup]

Returns x-y-line data grouped by the sets of "stable phases" (for example "LIQUID" or "LIQUID + FCC\_A1"). The available quantities can be found in the documentation of the factory class ThermodynamicQuantity.

**Note:** The different datasets might contain *NaN*-values between different subsections and different lines of an ambiguous dataset. They might not be sorted **even if the flag `sort\_and\_merge` has been set** (because they might be unsortable due to their nature).

**Note:** Its possible to use functions as axis variables, either by using ThermodynamicQuantity. user\_defined\_function, or by using an expression that contains '='.

Exampleget\_values\_grouped\_by\_quantity\_of('T',ThermodynamicQuantity.user\_defined\_function('HM.T'))

**Example** get\_values\_grouped\_by\_quantity\_of('T', 'CP=HM.T')

## Parameters

- **x\_quantity** The first quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example 'T'), *MATERIAL\_B\_FRACTION*, or even a function (for example 'f=T\*1.01')
- **y\_quantity** The second quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example '*NV*'), *MATERIAL\_B\_FRACTION*, or even a function (for example '*CP=HM.T*')
- **sort\_and\_merge** If *True*, the data will be sorted and merged into as few subsections as possible (divided by *NaN*)

#### Returns

Containing the datasets with the quantities as their keys

**get\_values\_of**(x\_quantity: Union[ThermodynamicQuantity, str], y\_quantity: Union[ThermodynamicQuantity, str])  $\rightarrow$  [List[float], List[float]]

Returns sorted x-y-line data without any separation. Use *get\_values\_grouped\_by\_quantity\_of()* or *get\_values\_grouped\_by\_stable\_phases\_of()* instead if you need such a separation. The available quantities can be found in the documentation of the factory class ThermodynamicQuantity.

**Note:** This method will always return sorted data without any *NaN*-values. If it is unsortable that might give data that is hard to interpret. In such a case you need to choose the quantity in another way or use one of the other methods. One example of this is to use quantities with *All*-markers, for example *MassFractionOfAComponent("All")*.

**Note:** Its possible to use functions as axis variables, either by using ThermodynamicQuantity. user\_defined\_function(), or by using an expression that contains '='.

**Example** get\_values\_grouped\_by\_quantity\_of('T', ThermodynamicQuantity.user\_defined\_function('HM.T'))

**Example** get\_values\_grouped\_by\_quantity\_of('T', 'CP=HM.T')

#### **Parameters**

- x\_quantity The first thermodynamic quantity ("x-axis"), Console Mode syntax strings can be used as an alternative (for example 'T', MATERIAL\_B\_FRACTION, or even a function (for example 'f=T\*1.01').
- **y\_quantity** The second thermodynamic quantity ("y-axis"), Console Mode syntax strings can be used as an alternative (for example '*NV*'), *MATERIAL\_B\_FRACTION*, or even a function (for example '*CP=HM.T*')

## Returns

A tuple containing the x- and y-data in lists

## save\_to\_disk(path: str)

Saves the result to disc. Note that a result is a folder, containing potentially many files. The result can later be loaded with load\_result\_from\_disk()

## **Parameters**

**path** – the path to the folder you want the result to be saved in. It can be relative or absolute.

## Returns

this MaterialToMaterialPropertyDiagramResult object

## set\_phase\_name\_style(phase\_name\_style\_enum: PhaseNameStyle = PhaseNameStyle.NONE)

Sets the style of the phase name labels that will be used in the result data object (constitution description, ordering description,  $\dots$ ).

**Default**: PhaseNameStyle.NONE

## Parameters

phase\_name\_style\_enum - The phase name style

## Returns

This MaterialToMaterialPropertyDiagramResult object

# class tc\_python.material\_to\_material.MaterialToMaterialSingleEquilibriumCalculation(calculator) Bases: AbstractSingleEquilibriumCalculation

Configuration for a Material to Material single fraction of B calculation.

**Note:** Specify the conditions and possibly other settings, the calculation is performed with *calculate()*.

## $calculate(timeout_in_minutes: float = 0.0) \rightarrow MaterialToMaterialSingleEquilibriumResult$

Performs the material to material calculation.

Note: The calculation result is no temporary result object.

#### **Parameters**

**timeout\_in\_minutes** – Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

#### Returns

A new *MaterialToMaterialSingleEquilibriumResult* object which can be used to get specific values from the calculated result. It is undefined behavior to use that object after the state of the calculation has been changed.

#### disable\_global\_minimization()

Turns the global minimization completely off.

## Returns

This MaterialToMaterialSingleEquilibriumCalculation object

## enable\_global\_minimization()

Turns the global minimization on (using the default settings).

## Returns

This MaterialToMaterialSingleEquilibriumCalculation object

#### $get\_components() \rightarrow List[str]$

Returns a list of components in the system (including all components auto-selected by the database(s)).

#### Returns

The components

## get\_gibbs\_energy\_addition\_for(phase: str) $\rightarrow$ float

Used to get the additional energy term (always being a constant) of a given phase. The value given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

#### Parameters

phase – Specify the name of the (stoichiometric or solution) phase with the addition

## Returns

Gibbs energy addition to G per mole formula unit.

## get\_system\_data() $\rightarrow$ SystemData

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using *with\_system\_modifications()*.

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as \*.tdb-file.

#### Returns

The system data

#### run\_poly\_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine.

## Parameters command – The Thermo-Calc Console Mode command

#### Returns

This MaterialToMaterialSingleEquilibriumCalculation object

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

## set\_activities(activities: Dict[str, float])

Sets the constant activity conditions.

Note: The activity conditions are identical for both materials.

#### **Parameters**

activities – The constant activities

#### Returns

This MaterialToMaterialSingleEquilibriumCalculation object

## set\_component\_to\_entered(component: str)

Sets the specified component to the status ENTERED, that is the default state.

## Parameters

component – The component name or ALL\_COMPONENTS

#### Returns

This MaterialToMaterialSingleEquilibriumCalculation object

## set\_component\_to\_suspended(component: str, reset\_conditions: bool = False)

Sets the specified component to the status SUSPENDED, i.e. it is ignored in the calculation.

#### **Parameters**

• **reset\_conditions** – if 'True' also remove composition conditions for the component if they are defined

• **component** – The component name or *ALL\_COMPONENTS* 

## Returns

This MaterialToMaterialSingleEquilibriumCalculation object

set\_composition\_unit(unit: CompositionUnit = CompositionUnit.MASS\_PERCENT)

Sets the composition unit of both materials A and B.

**Default**: Weight percent

#### **Parameters**

unit – The composition unit of both materials A and B

#### Returns

This MaterialToMaterialSingleEquilibriumCalculation object

## set\_gibbs\_energy\_addition\_for(phase: str, gibbs\_energy: float)

Used to specify the additional energy term (always being a constant) of a given phase. The value (*gibbs\_energy*) given is added to the Gibbs energy of the (stoichiometric or solution) phase. It can represent a nucleation barrier, surface tension, elastic energy, etc.

It is not composition-, temperature- or pressure-dependent.

## Parameters

- phase Specify the name of the (stoichiometric or solution) phase with the addition
- gibbs\_energy Addition to G per mole formula unit

#### Returns

This MaterialToMaterialSingleEquilibriumCalculation object

**set\_material\_a**(composition: Dict[str, float], dependent\_component: Optional[str] = None)

Sets the composition of the material A.

The unit is set with set\_composition\_unit().

**Tip:** The material can also have constant activity conditions, they are set in *set\_activities(*).

## **Parameters**

- composition The composition of the material A
- **dependent\_component** The dependent component of the material A

#### Returns

This MaterialToMaterialSingleEquilibriumCalculation object

set\_material\_b(composition: Dict[str, float], dependent\_component: Optional[str] = None)
Sets the composition of the material B.

The unit is set with set\_composition\_unit().

**Tip:** The material can also have constant activity conditions, they are set in set\_activities().

#### **Parameters**

• composition – The composition of the material B

• dependent\_component – The dependent component of the material B

#### Returns

This MaterialToMaterialSingleEquilibriumCalculation object

## set\_phase\_to\_dormant(phase: str)

Sets the phase to the status DORMANT, necessary for calculating the driving force to form the specified phase.

## Parameters

phase – The phase name or ALL\_PHASES for all phases

#### Returns

This MaterialToMaterialSingleEquilibriumCalculation object

#### set\_phase\_to\_entered(phase: str, amount: float = 1.0)

Sets the phase to the status ENTERED, that is the default state.

## **Parameters**

• phase – The phase name or ALL\_PHASES for all phases

• **amount** – The phase fraction (between 0.0 and 1.0)

#### Returns

This MaterialToMaterialSingleEquilibriumCalculation object

#### set\_phase\_to\_fixed(phase: str, amount: float)

Sets the phase to the status FIXED, i.e. it is guaranteed to have the specified phase fraction after the calculation.

#### **Parameters**

• phase – The phase name

• **amount** – The fixed phase fraction (between 0.0 and 1.0)

#### Returns

This MaterialToMaterialSingleEquilibriumCalculation object

#### set\_phase\_to\_suspended(phase: str)

Sets the phase to the status SUSPENDED, i.e. it is ignored in the calculation.

### Parameters

**phase** – The phase name or *ALL\_PHASES* for all phases

#### Returns

This MaterialToMaterialSingleEquilibriumCalculation object

#### set\_pressure(pressure: float)

Sets the pressure (i.e. the condition P).

**Note:** If the flag *default\_conditions=True* has been set during the creation of the calculator, the pressure is set to 1000 hPa by default.

#### **Parameters**

pressure – The pressure [Pa]

#### Returns

This MaterialToMaterialSingleEquilibriumCalculation object

#### set\_system\_size(system\_size: float)

Sets the system size (i.e. the condition 'N', the number of moles).

**Note:** If the flag *default\_conditions=True* has been set during the creation of the calculator, the system size is set to 1.0 moles by default.

#### Parameters

**system\_size** – The system size [mole]

#### Returns

This MaterialToMaterialSingleEquilibriumCalculation object

#### with\_first\_constant\_condition(condition: ConstantCondition)

Sets the first constant condition (either temperature of fraction of material B).

## Parameters

condition – The condition

#### Returns

This MaterialToMaterialSingleEquilibriumCalculation object

## with\_options(options: SingleEquilibriumOptions)

Sets the simulation options.

**Parameters options** – The simulation options

#### Returns

This SingleEquilibriumCalculation object

## with\_reference\_state(component: str, phase: str = 'SER', temperature: float = -1.0, pressure: float = 100000.0)

The reference state for a component is important when calculating activities, chemical potentials and enthalpies and is determined by the database being used. For each component the data must be referred to a selected phase, temperature and pressure, i.e. the reference state.

All data in all phases where this component dissolves must use the same reference state. However, different databases can use different reference states for the same element/component. It is important to be careful when combining data obtained from different databases.

By default, activities, chemical potentials and so forth are computed relative to the reference state used by the database. If the reference state in the database is not suitable for your purposes, use this command to set the reference state for a component using SER, i.e. the Stable Element Reference (which is usually set as default for a major component in alloys dominated by the component). In such cases, the temperature and pressure for the reference state is not needed.

For a phase to be usable as a reference for a component, the component needs to have the same composition as an end member of the phase. The reference state is an end member of a phase. The selection of the end member associated with the reference state is only performed once this command is executed.

If a component has the same composition as several end members of the chosen reference phase, then the end member that is selected at the specified temperature and pressure will have the lowest Gibbs energy.

#### Parameters

- component The name of the element must be given.
- **phase** Name of a phase used as the new reference state. Or SER for the Stable Element Reference.
- **temperature** The Temperature (in K) for the reference state. Or CURRENT\_TEMPERATURE which means that the current temperature is used at the time of evaluation of the reference energy for the calculation.
- **pressure** The Pressure (in Pa) for the reference state.

#### Returns

This MaterialToMaterialSingleEquilibriumCalculation object

#### with\_second\_constant\_condition(condition: ConstantCondition)

Sets the second constant condition (either temperature of fraction of material B).

Parameters condition – The condition

Returns

This MaterialToMaterialSingleEquilibriumCalculation object

#### with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

**Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

#### **Parameters**

system\_modifications – The system modification to be performed

#### Returns

This MaterialToMaterialSingleEquilibriumCalculation object

# class tc\_python.material\_to\_material.MaterialToMaterialSingleEquilibriumResult(result)

Bases: SingleEquilibriumResult

Result of a Material To Material calculation for a single fraction of material B, it can be evaluated using a quantity or Console Mode syntax.

## change\_pressure(pressure: float)

Change the pressure and re-evaluate the results from the equilibrium without minimizing Gibbs energy, i.e. with higher performance. The properties are calculated at the new pressure using the phase amount, temperature and composition of phases from the initial equilibrium. Use *get\_value\_of()* to obtain them.

#### Parameters

**pressure** – The pressure [Pa]

Returns

This MaterialToMaterialSingleEquilibriumResult object

# change\_temperature(temperature: float)

Change the temperature and re-evaluate the results from the equilibrium without minimizing Gibbs energy, i.e. with high performance. The properties are calculated at the new temperature using the phase amount, pressure and composition of phases from the initial equilibrium. Use *get\_value\_of()* to obtain them.

**Note:** This is typically used when calculating room temperature properties (e.g. density) for a material when it is assumed that the equilibrium phase amount and composition freeze-in at a higher temperature during cooling.

## Parameters temperature – The temperature [K]

#### Returns

This MaterialToMaterialSingleEquilibriumResult object

## $\texttt{get\_components()} \rightarrow List[str]$

Returns the names of the components selected in the system (including any components auto-selected by the database(s)).

#### Returns

The names of the selected components

# $\texttt{get\_conditions()} \rightarrow List[str]$

Returns the conditions.

# Returns

The selected conditions

#### $get_phases() \rightarrow List[str]$

Returns the phases present in the system due to its configuration. It also contains all phases that have been automatically added during the calculation, this is the difference to the method System. get\_phases\_in\_system().

#### Returns

The names of the phases in the system including automatically added phases

# $\texttt{get\_stable\_phases()} \rightarrow List[str]$

Returns the stable phases (i.e. the phases present in the current equilibrium).

#### Returns

The names of the stable phases

## **get\_value\_of**(*quantity: Union*[ThermodynamicQuantity, *str*]) → float

Returns a value from a single equilibrium calculation.

# Parameters

**quantity** – The thermodynamic quantity to get the value of; a Console Mode syntax strings can be used as an alternative (for example "NPM(FCC\_A1)")

## Returns

The requested value

# run\_poly\_command(command: str)

Runs a Thermo-Calc command from the Console Mode POLY module immediately in the engine. This affects only the state of the result object.

# Parameters

**command** – The Thermo-Calc Console Mode command

Returns

This MaterialToMaterialSingleEquilibriumResult object

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

save\_to\_disk(path: str)

Saves the result to disk. Note that the result is a folder, containing potentially many files. The result can later be loaded with load\_result\_from\_disk()

# **Parameters**

path – the path to the folder you want the result to be saved in. It can be relative or absolute.

Returns

this MaterialToMaterialSingleEquilibriumResult object

*start\_temperature: float = 2000*)

# Bases: MaterialToMaterialCalculationAxis

A temperature calculation axis.

```
class tc_python.material_to_material.TemperatureCondition(temperature: float = 1000.0)
```

Bases: ConstantCondition

A constant temperature condition.

# 6.1.9 Package "process\_metallurgy"

# 6.1.9.1 Module "base"

# class tc\_python.process\_metallurgy.base.AbstractAddition

Bases: object

The base class for representing an addition to an equilibrium calculation or process simulation.

# $\texttt{get\_composition()} \rightarrow Dict[str, float]$

Returns the composition of the addition - without containing a dependent component.

# Returns

The composition [in the unit provided by getCompositionUnit()]

## abstract get\_composition\_unit()

Returns the composition unit used in this addition.

# Returns

The composition unit

# get\_dependent\_component() $\rightarrow$ str

Returns the dependent component.

#### Returns

The dependent component or an empty string if no dependent component is defined

# $get_elements() \rightarrow Set[str]$

Returns all elements of the addition.

Returns

The elements

# $\texttt{get\_id}() \to str$

Returns the unique ID of the addition.

## Returns

The unique ID of the addition

# get\_temperature() $\rightarrow$ float

Returns the temperature of the addition. This refers to the temperature before it is added to the process.

#### Returns

The temperature [K]

# $\texttt{is\_do\_scale()} \rightarrow \texttt{bool}$

Returns if the composition of the addition is being scaled to 100% / 1 or not.

#### Returns

If the composition is scaled

# $\texttt{is\_empty()} \rightarrow \texttt{bool}$

Returns if the addition is "empty", i.e., has zero amount.

# Returns

If the addition is empty

# class tc\_python.process\_metallurgy.base.ActivityReference(value)

Bases: Enum

The reference for a slag activity calculation. The actual reference phase depends on the component for which the activity is request and can be obtained by using these methods on the result object:  $tc_python.process_metallurgy.process.ProcessSimulationResult.get_formula_for_activity_of_slag() or <math>tc_python.process_metallurgy.equilibrium.$ EquilibriumResult.get\_formula\_for\_activity\_of\_slag().

# LIQUID = 1

The reference is liquid slag.

# SOLID = 0

The reference is solid slag.

# class tc\_python.process\_metallurgy.base.PhaseGroup(value)

# Bases: Enum

The phase group, such a group is collecting all phases that belong to a certain type.

# $ALL_METAL = 2$

All metal phases.

# $ALL_SLAG = 5$

All slag phases.

# GAS = 6

All gas phases.

#### $LIQUID_METAL = 0$

All liquid metal phases.

# $LIQUID_SLAG = 4$

All liquid slag phases.

# $SOLID_METAL = 1$

All solid metal phases.

# $SOLID_SLAG = 3$

All solid slag phases.

# class tc\_python.process\_metallurgy.base.ProcessDatabase(value)

# Bases: Enum

The database used for a Process Metallurgy calculation.

# LATEST = 0

The latest database available.

# OXDEMO = 1

The database OXDEMO.

# TCOX10 = 4

The database TCOX10.

# TCOX11 = 5

The database TCOX11.

# TCOX12 = 6

The database TCOX12.

# TCOX13 = 7

The database TCOX13.

# TCOX14 = 8

The database TCOX14.

# TCOX8 = 2

The database TCOX8.

# TCOX9 = 3

The database TCOX9.

# $\texttt{get\_name()} \to \texttt{str}$

Returns the name of the actually used database.

Tip: This is especially useful if LATEST is used.

# Returns

The name of the database

# class tc\_python.process\_metallurgy.base.ProcessMetallurgyOptions Bases: SingleEquilibriumOptions

The options for a process metallurgy calculation.

# disable\_approximate\_driving\_force\_for\_metastable\_phases()

Disables the approximation of the driving force for metastable phases.

Default: Enabled

**Note:** When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it is important that these driving forces are correct, use *disable\_approximate\_driving\_force\_for\_metastable\_pha* to force the calculation to converge for the metastable phases.

#### Returns

This ProcessMetallurgyOptions object

# disable\_control\_step\_size\_during\_minimization()

Disables step size control during minimization (non-global).

**Default**: Enabled

#### Returns

This ProcessMetallurgyOptions object

# disable\_force\_positive\_definite\_phase\_hessian()

Disables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

Default: Enabled

#### Returns

This ProcessMetallurgyOptions object

# enable\_approximate\_driving\_force\_for\_metastable\_phases()

Enables the approximation of the driving force for metastable phases.

Default: Enabled

**Note:** When enabled, the metastable phases are included in all iterations. However, these may not have reached their most favorable composition and thus their driving forces may be only approximate.

If it is important that these driving forces are correct, use *disable\_approximate\_driving\_force\_for\_metastable\_pha* to force the calculation to converge for the metastable phases.

#### Returns

This ProcessMetallurgyOptions object

# enable\_control\_step\_size\_during\_minimization()

Enables step size control during normal minimization (non-global).

Default: Enabled

# Returns

This ProcessMetallurgyOptions object

#### enable\_force\_positive\_definite\_phase\_hessian()

Enables forcing of positive definite phase Hessian. This determines how the minimum of an equilibrium state in a normal minimization procedure (non-global) is reached. For details, search the Thermo-Calc documentation for "Hessian minimization".

## Default: Enabled

#### Returns

This ProcessMetallurgyOptions object

#### set\_global\_minimization\_max\_grid\_points(max\_grid\_points: int = 2000)

Sets the maximum number of grid points in global minimization. **Only applicable if global minimization is actually used**.

Default: 2000 points

Parameters max\_grid\_points – The maximum number of grid points

Returns This ProcessMetallurgyOptions object

set\_max\_no\_of\_iterations(max\_no\_of\_iterations: int = 2000)

Sets the maximum number of iterations for the CALPHAD minimizer.

Default: max. 2000 iterations

**Note:** As some models give computation times of more than 1 CPU second/iteration, this number is also used to check the CPU time and the calculation stops if 500 CPU seconds/iterations are used.

#### **Parameters**

max\_no\_of\_iterations – The max. number of iterations

#### Returns

This ProcessMetallurgyOptions object

# set\_process\_minimization\_policy(minimization\_policy: ProcessMinimizationPolicy)

Sets the minimization policy for the process metallurgy calculation. It is possible to choose policies that try different methods if one method fails.

## **Parameters**

**minimization\_policy** – The minimization policy to be used

Returns

This ProcessMetallurgyOptions object

#### set\_required\_accuracy(accuracy: float = 1e-06)

Sets the required relative accuracy.

Default: 1.0E-6

**Note:** This is a relative accuracy, and the program requires that the relative difference in each variable must be lower than this value before it has converged. A larger value normally means fewer iterations but less accurate solutions. The value should be at least one order of magnitude larger than the machine precision.

#### Parameters

accuracy – The required relative accuracy

## Returns

This ProcessMetallurgyOptions object

# set\_smallest\_fraction(smallest\_fraction: float = 1e-16)

Sets the smallest fraction for constituents that are unstable.

It is normally only in the gas phase that you can find such low fractions.

The **default value** for the smallest site-fractions is 1E-16 for all phases except for IDEAL phase with one sublattice site (such as the GAS mixture phase in many databases) for which the default value is always as 1E-30.

# Parameters smallest\_fraction – The smallest fraction for constituents that are unstable

Returns

This ProcessMetallurgyOptions object

class tc\_python.process\_metallurgy.base.ProcessMinimizationPolicy(value)

# Bases: Enum

The policy for the CALPHAD minimization routine used in a calculation.

**Note:** This affects the runtime and stability of a calculation. Global minimization is more stable but more timeconsuming. Local minimization is much faster but can miss new phases coming up. Global test is a compromise between both approaches.

# GLOBAL = 0

Always use global minimization.

#### $GLOBAL_TEST = 1$

Always use global test.

# GLOBAL\_TEST\_\_GLOBAL = 5

First try global test, if that fails use global minimization.

# LOCAL = 2

Always use local minimization.

## $LOCAL\_GLOBAL\_TEST = 3$

First try local minimization, if that fails use global test.

# LOCAL\_\_GLOBAL\_TEST\_\_GLOBAL = 4

First try local minimization, if that fails try global test and if that fails use global minimization.

## class tc\_python.process\_metallurgy.base.SlagProperty(value)

# Bases: Enum

The slag property, different definitions are available. The actual definition of a certain slag property for the current system can be obtained using these methods on the result object: tc\_python.process\_metallurgy.process.ProcessSimulationResult.getFormulaForSlagProperty() or tc\_python.process\_metallurgy.equilibrium.EquilibriumResult.getFormulaForSlagProperty().

**Note:** If not all components required by the definition of slag property are available in a given system, the slag property will return *NaN*.

# B2 = 0

Basicity B2 (based on: CaO / SiO2).

# B3 = 1

Basicity B3 (based on: CaO / MgO / SiO2).

# B4 = 3

Basicity B4 (based on: CaO / MgO / SiO2 / Al2O3).

# BAS2 = 2

Basicity Bas2 (based on: CaO / MgO / SiO2 / Al2O3 / TiO2).

# $BELLS_RATIO = 4$

Basicity Bell's ratio (based on: CaO / MgO / SiO2 / Al2O3).

# LOG\_10\_SULPHUR\_CAPACITY = 5

Logarithmic sulphur capacity of the slag.

# LS = 6

Sulphur capacity *Ls* of the slag.

class tc\_python.process\_metallurgy.base.SlagType(value)

# Bases: Enum

The type of slag considered for a slag property calculation.

# ALL = 2

The slag property is calculated for all slag, i.e. for both the liquid and solid slag.

# LIQUID = 1

The slag property is calculated for all liquid slag.

# SOLID = 0

The slag property is calculated for all solid slag.

# 6.1.9.2 Module "equilibrium"

class tc\_python.process\_metallurgy.equilibrium.AbstractEquilibriumAddition
 Bases: AbstractAddition

The base class for representing an addition to an equilibrium calculation.

#### set\_amount(amount: float)

Change the amount of the addition.

# Parameters

amount – The new amount [in the amount unit of this addition]

# Returns

This AbstractEquilibriumAddition object

# set\_component\_composition(component: str, content: float)

Change the composition of a component of the addition.

# Parameters

- component The component to be changed
- **content** The new content of the component [in the composition unit defined for this addition]

#### Returns

This AbstractEquilibriumAddition object

# class tc\_python.process\_metallurgy.equilibrium.AdiabaticEquilibriumCalculation(calculator) Bases: EquilibriumCalculation

An *adiabatic* Process Metallurgy equilibrium calculation. Such calculations can for example be used to determine the global equilibrium state of a process.

#### add\_addition(addition: AbstractEquilibriumAddition)

Adds an addition to the calculation.

#### Parameters

addition - A EquilibriumAddition or EquilibriumGasAddition

#### Returns

This AdiabaticEquilibriumCalculation object

#### add\_poly\_command(command: str)

Adds a Thermo-Calc Console syntax POLY module command which will be executed when performing the calculate() method.

If multiple commands are added, they will be executed in the order of addition. Each command will only be executed one.

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

#### **Parameters**

command – The POLY module command in Thermo-Calc console syntax

## Returns

This AdiabaticEquilibriumCalculation object

# $calculate(timeout_in_minutes: float = 0.0) \rightarrow EquilibriumResult$

Runs the Process Metallurgy equilibrium calculation.

#### **Parameters**

timeout\_in\_minutes – The calculation will be aborted after that time, default: no timeout

#### Returns

A new EquilibriumResult object

#### remove\_addition(addition: AbstractEquilibriumAddition)

Removes an addition from the calculation.

# Parameters

**addition** – The addition to be removed

# Returns

This AdiabaticEquilibriumCalculation object

# remove\_all\_additions()

Removes all additions from the calculation.

# Returns

This AdiabaticEquilibriumCalculation object

set\_pressure(pressure: float = 100000.0)

Sets the pressure.

Parameters pressure – The pressure [Pa]

Returns

This AdiabaticEquilibriumCalculation object

with\_options(options: ProcessMetallurgyOptions)

Sets the options for the calculation.

**Parameters options** – The options

#### Returns

This AdiabaticEquilibriumCalculation object

class tc\_python.process\_metallurgy.equilibrium.EquilibriumAddition(composition: Dict[str, float],

amount: float, temperature: float = 293.15, composition\_unit: CompositionUnit = CompositionUnit.MASS\_PERCENT, do\_scale: bool = False)

Bases: AbstractEquilibriumAddition

An addition to an equilibrium calculation.

**Tip:** By setting  $do\_scale=True$ , the composition will be scaled to 100% / fraction of 1. This is useful if the composition provided is not summing to 100% / 1. An example could be a slag addition which is provided like this: 90 wt-% CaO - 5 wt-% Al2O3 - 4 wt-% SiO2.

# Parameters

- composition The composition
- amount The amount [kg]
- temperature The initial addition temperature (default: 20 °C) [K]
- composition\_unit The composition unit
- **do\_scale** If the composition is scaled to 100% / fraction of 1

# $\texttt{get\_amount()} \to \texttt{float}$

Returns the amount of this addition.

## Returns

The amount [kg]

# $\texttt{get\_composition\_unit()} \rightarrow \textit{CompositionUnit}$

Returns the composition unit used in this addition.

#### Returns

The composition unit

class tc\_python.process\_metallurgy.equilibrium.EquilibriumCalculation(metallurgical\_reaction)
 Bases: object

A Process Metallurgy equilibrium calculation. Such calculations can for example be used to determine the global equilibrium state of a process.

abstract add\_addition(addition: AbstractEquilibriumAddition)

Adds an addition to the calculation.

Parameters addition – The addition

## Returns

This EquilibriumCalculation object

# abstract add\_poly\_command(command: str)

Adds a Thermo-Calc Console syntax POLY module command which will be executed when performing the calculate() method.

If multiple commands are added, they will be executed in the order of addition. Each command will only be executed one.

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

#### **Parameters**

command – The POLY module command in Thermo-Calc console syntax

## Returns

This EquilibriumCalculation object

**abstract calculate**(*timeout\_in\_minutes: float* = 0.0)  $\rightarrow$  *EquilibriumResult* 

Runs the Process Metallurgy equilibrium calculation.

#### **Parameters**

timeout\_in\_minutes – The calculation will be aborted after that time, default: no timeout

#### Returns

A new EquilibriumResult object

abstract remove\_addition(addition: AbstractEquilibriumAddition)

Removes an addition from the calculation.

# Parameters

**addition** – The addition to be removed

## Returns

This EquilibriumCalculation object

# abstract remove\_all\_additions()

Removes all additions from the calculation.

## Returns

This EquilibriumCalculation object

**abstract set\_pressure**(*pressure: float* = 100000.0)

Sets the pressure.

**Parameters pressure** – The pressure [Pa]

Returns

This EquilibriumCalculation object

## update\_addition(addition: AbstractEquilibriumAddition)

Replaces an already added addition with an updated one. This is usually used to change the composition or amount of an addition while iterating over them. Typically, this is done for stepping or mapping calculations.

Note: The calculation must already contain the addition object to be updated.

# Parameters

addition – The new addition containing updated values

Returns

This IsoThermalMetallurgyCalculation object

# abstract with\_options(options: ProcessMetallurgyOptions)

Sets the options for the calculation.

Parameters options – The options

# Returns

This EquilibriumCalculation object

class tc\_python.process\_metallurgy.equilibrium.EquilibriumGasAddition(composition: Dict[str,

float], amount: float, temperature: float = 293.15, amount\_unit: GasAmountUnit.NORM\_CUBIC\_METER, composition\_unit: GasCompositionUnit.VOLUME\_PERCENT, do\_scale: bool = False)

Bases: AbstractEquilibriumAddition

A gas addition to an equilibrium calculation.

**Tip:** By setting *do\_scale=True*, the composition will be scaled to 100% / fraction of 1. This is useful if the composition provided is not summing to 100% / 1. An example could be a gas addition which is provided like this: 90 vol-% Ar - 10 vol-% O2.

 $\texttt{get\_amount()} \to \texttt{float}$ 

Returns the amount of this addition.

**Note:** The amount unit can be obtained using *get\_amount\_unit(*).

#### Returns

The amount [in the amount unit]

 $get\_amount\_unit() \rightarrow GasAmountUnit$ 

Returns the amount unit used in this addition.

Returns

The amount unit

# $get\_composition\_unit() \rightarrow GasCompositionUnit$

Returns the composition unit used in this addition.

## Returns

The composition unit

class tc\_python.process\_metallurgy.equilibrium.EquilibriumResult(result)

Bases: AbstractResult

The result of a Process Metallurgy equilibrium calculation.

# $\texttt{get\_activity\_of\_slag}(\textit{component: str, reference: ActivityReference} = \textit{ActivityReference.LIQUID}) \rightarrow \texttt{activityReference} = \texttt{Activ$

float

Returns the activity of a component in the slag.

#### Parameters

- **component** The component
- reference The reference for the activity, can be liquid or solid slag, default: liquid slag

#### Returns

The activity of the component [-]

# $\texttt{get\_amount()} \to \texttt{float}$

Returns the total amount.

# Returns

The total amount [kg]

# $\texttt{get\_amount\_of\_elements()} \rightarrow Dict[str, float]$

Returns the amount of each element.

# Returns

The amount of the elements [kg]

# $\texttt{get\_amount\_of\_phase\_groups()} \rightarrow \texttt{Dict}[\textit{PhaseGroup}, \texttt{float}]$

Returns the amount of each phase group (e.g., for example all liquid slag).

#### Returns

The amount of the phase groups [kg]

# $\texttt{get\_amount\_of\_phases()} \rightarrow Dict[str, float]$

Returns the amount of each phase.

# Returns

The amount of the phases [kg]

 $get\_components() \rightarrow Set[str]$ 

Returns all components defined for the elements present in this result.

#### Returns

The components present in this result

 $\texttt{get\_composition}(\textit{composition\_unit: CompositionUnit} = \textit{CompositionUnit.MASS\_PERCENT}) \rightarrow \texttt{Dict[str},$ 

float]

Returns the composition of the result.

## Parameters

composition\_unit - The composition unit, default: mass percent

#### Returns

The composition

get\_composition\_of\_phase(phase: str, composition\_unit: CompositionUnit =

*CompositionUnit.MASS\_PERCENT, composition\_type:* CompositionType = *CompositionType.COMPONENT*) → Dict[str, float]

Returns the composition of a phase in the result.

## Parameters

- **phase** The phase name
- composition\_unit The composition unit, default: mass percent
- **composition\_type** Defines if the composition is given by element (e.g., 75 wt-% Fe 25 wt-% Cr) or by component (e.g. 65 wt-% Al2O3 35 wt-% CaO). In case of a metallic phase, the composition is given by element even if *component* is selected. **Default: by component**.

# Returns

The composition

 $\label{eq:get_composition_of_phase_group} (phase_group: PhaseGroup, composition_unit: CompositionUnit = CompositionUnit.MASS_PERCENT, composition_type: CompositionType = CompositionType.COMPONENT) \rightarrow Dict[str, float]$ 

Returns the composition of a phase group (e.g., all liquid slag) in the result.

# Parameters

- phase\_group The phase group
- composition\_unit The composition unit, default: mass percent
- **composition\_type** Defines if the composition is given by element (e.g., 75 wt-% Fe 25 wt-% Cr) or by component (e.g. 65 wt-% Al2O3 35 wt-% CaO). In case of a metallic phase, the composition is given by element even if *component* is selected. **Default: by component**.

#### Returns

The composition

# $\texttt{get\_elements()} \rightarrow Set[str]$

Returns all elements defined for the result.

# Returns

All elements present in this result

# $get\_formula\_for\_activity\_of\_slag(component: str, reference: ActivityReference = ActivityReference.LIQUID) \rightarrow str$

Returns the Thermo-Calc Console syntax formula used for calculating the activity of a component in the slag (e.g. *AC(AL2O3, IONIC\_LIQ)*. The actual activity can be obtained using *get\_activity\_of\_slag()*.

#### **Parameters**

- component The component
- **reference** The reference for the activity, can be liquid or solid slag, **default: liquid slag**

#### Returns

The formula for calculating the activity

```
get_formula_for_slag_property(slag_property: SlagProperty, slag_type: SlagType=SlagType.ALL) → str
```

Returns the Thermo-Calc Console syntax formula used for calculating a property of the slag (e.g. B(CAO)/B(SIO2)). The actual slag property can be obtained using  $get_slag_property()$ .

#### **Parameters**

- slag\_property The slag property
- **slag\_type** The part of the slag for which the property will be calculated. Can be all slag, the liquid or the solid slag. **Default: all slag**

#### Returns

The formula for calculating the slag property

 $get\_fraction\_of\_phase\_groups(unit: PhaseUnit=PhaseUnit.MASS\_FRACTION) \rightarrow Dict[PhaseGroup, float]$ 

Returns the fraction of the phase groups (e.g., all liquid slag) in the result.

# Parameters

unit - The unit of the fraction, default: volume fraction

# Returns

The phase fractions

**get\_fraction\_of\_phases**(*unit:* PhaseUnit = PhaseUnit.MASS\_FRACTION)  $\rightarrow$  Dict[str, float]

Returns the fraction of the stable phases in the result.

#### **Parameters**

unit – The unit of the fraction, default: volume fraction

## Returns

The phase fractions

# $\texttt{get\_gas\_components()} \rightarrow \texttt{Set[str]}$

Returns all components of the gas phase defined for the elements present in this result.

# Returns

The components of the gas phase present in this result

# get\_oxygen\_partial\_pressure() $\rightarrow$ float

Returns the partial pressure of oxygen in the result.

# Returns

The partial pressure [Pa]

## get\_pressure() $\rightarrow$ float

Returns the pressure in the result.

#### Returns

The pressure [Pa]

**get\_slag\_property**(*slag\_property*: SlagProperty, *slag\_type*: SlagType = *SlagType*.*ALL*)  $\rightarrow$  float

Returns a property of the slag. These properties are mostly used to describe the property of a slag to pick up sulfur.

## **Parameters**

- slag\_property The slag property
- **slag\_type** The part of the slag for which the property will be calculated. Can be all slag, the liquid or the solid slag. **Default: all slag**

#### Returns

The slag property [unit depending on the property]

#### get\_stable\_phases() $\rightarrow$ Set[str]

Returns the stable phases in the result.

# Returns

The stable phases

# $get\_stable\_phases\_in\_phase\_group(phase\_group: PhaseGroup) \rightarrow Set[str]$

Returns the stable phases of a phase group (e.g., all liquid slag) in the result.

# Parameters phase\_group – The phase group

# Returns

The stable phases

# get\_temperature() $\rightarrow$ float

Returns the temperature in the result.

# Returns

The temperature [K]

# $get_value_of(classic_expression: str) \rightarrow float$

Returns a value for a thermodynamic quantity.

**Warning:** It should normally not be required to use this method, use the appropriate method available in the API instead.

## Parameters

**classic\_expression** – The thermodynamic quantity to get the value of in Thermo-Calc Console Mode syntax (for example "NPM(FCC\_A1)")

#### Returns

The requested value

# get\_viscosity\_dynamic\_of\_phase(phase: str) → float

Returns the dynamic viscosity of a phase in the result.

# Parameters

phase – The phase name

Returns

The dynamic viscosity [Pa\*s]

# get\_viscosity\_kinematic\_of\_phase(phase: str) → float

Returns the kinematic viscosity of a phase in the result.

Parameters phase – The phase name

Returns

The kinematic viscosity [m\*\*2/s]

class tc\_python.process\_metallurgy.equilibrium.IsoThermalEquilibriumCalculation(calculation)
Bases: EquilibriumCalculation

An *isothermal* Process Metallurgy equilibrium calculation. Such calculations can for example be used to determine the global equilibrium state of a process.

add\_addition(addition: AbstractEquilibriumAddition)

Adds an addition to the calculation.

```
Parameters
addition – A EquilibriumAddition or EquilibriumGasAddition
```

Returns This IsoThermalEquilibriumCalculation object

#### add\_poly\_command(command: str)

Adds a Thermo-Calc Console syntax POLY module command which will be executed when performing the calculate() method.

If multiple commands are added, they will be executed in the order of addition. Each command will only be executed one.

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw Thermo-Calc commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten equals sign).

## Parameters

command – The POLY module command in Thermo-Calc console syntax

#### Returns

This IsoThermalEquilibriumCalculation object

 $calculate(timeout_in_minutes: float = 0.0) \rightarrow EquilibriumResult$ 

Runs the Process Metallurgy equilibrium calculation.

## Parameters

timeout\_in\_minutes - The calculation will be aborted after that time, default: no timeout

#### Returns

A new EquilibriumResult object

# remove\_addition(addition: AbstractEquilibriumAddition)

Removes an addition from the calculation.

# Parameters

addition – The addition to be removed

### Returns

This IsoThermalEquilibriumCalculation object

# remove\_all\_additions()

Removes all additions from the calculation.

## Returns

This IsoThermalEquilibriumCalculation object

#### set\_pressure(pressure: float = 100000.0)

Sets the pressure.

# Parameters

pressure – The pressure [Pa]

# Returns

This IsoThermalEquilibriumCalculation object

# set\_temperature(temperature: float)

Sets the temperature.

Parameters temperature – The temperature [K]

#### Returns

This IsoThermalEquilibriumCalculation object

#### update\_addition(addition: AbstractEquilibriumAddition)

Replaces an already added addition with an updated one.

**Tip:** This is usually used to change the composition or amount of an addition while iterating over multiple values. Typically, this is done for stepping or mapping calculations.

Note: The calculation must already contain the addition object to be updated.

#### Parameters

addition - A previously added addition object with the updated values

#### Returns

This IsoThermalEquilibriumCalculation object

#### with\_options(options: ProcessMetallurgyOptions)

Sets the options for the calculation.

# Parameters

options – The options

# Returns

This IsoThermalEquilibriumCalculation object

# 6.1.9.3 Module "process"

# class tc\_python.process\_metallurgy.process.AbstractContinuousAddition Bases: AbstractProcessAddition

The base class representing an addition in a process simulation that is added continuously over a period of time.

# class tc\_python.process\_metallurgy.process.AbstractProcessAddition

Bases: AbstractAddition

The base class for representing an addition in a process simulation.

#### class tc\_python.process\_metallurgy.process.AbstractSingleTimeAddition

Bases: AbstractProcessAddition

The base class representing an addition in a process simulation that is added at a distinct time point.

Bases: Zone

A bulk zone in a process simulation, this is representing a large volume in the process, for example the steel melt or the top slag. A zone is a volume in a process that has identical temperature and composition. It has well-defined boundaries to other zones.

Tip: This is a generic class and seldom used directly. Use instead MetalBulkZone or SlagBulkZone.

## **add\_addition**(*addition*: AbstractSingleTimeAddition, *time: float = 0.0*)

Adds a single-time addition at the specified time point to the zone. The addition will be dissolved immediately.

# Parameters

- addition A SingleTimeAddition or SingleTimeGasAddition
- time The time point [s]

#### Returns

This BulkZone object

Adds a constant addition continuously during the specified time period to the zone. All added material will be dissolved immediately.

# Parameters

- addition A ContinuousAddition or ContinuousGasAddition
- from\_time The start time point [s]
- **to\_time** The end time point [s]

# Returns

This BulkZone object

**add\_power**(*power: float, from\_time: float = 0.0, to\_time: float = nan*)

Adds a constant power during a specified time period to the zone (for example heating or cooling).

#### **Parameters**

- power The power [W]
- from\_time The start time point [s]
- **to\_time** The end time point [s]

## Returns

This BulkZone object

# disable\_degassing()

Disables degassing for this zone, i.e. all gas formed at any time step will be staying in this zone.

#### Returns

This BulkZone object

# enable\_degassing()

Enables degassing for this zone, i.e. any gas formed at any time step will be removed after that time step. This gas will be transferred into the *exhaust gas zone*. **This is the default**.

#### Returns

This BulkZone object

# get\_density() $\rightarrow$ float

Returns the density of the zone

Returns

The density [kg/m\*\*3]

# $\texttt{get\_elements()} \rightarrow Set[str]$

Returns the elements present in the zone. The elements are determined by the additions.

# Returns

The elements

# $\texttt{get\_id}() \to \text{str}$

Returns the unique id of the zone. :return: The zone id

## get\_phase\_group\_to\_transfer() $\rightarrow$ *PhaseGroup*

Returns the phase group that is transferred from the attached reaction zones back to this zone after each time step.

#### Returns

The phase group

# $\texttt{is\_degassing\_enabled()} \rightarrow \texttt{bool}$

Returns if degassing is enabled in the zone.

## Returns

If degassing is enabled

Bases: AbstractContinuousAddition

An addition in a process simulation that is added continuously during a period of time.

It is assumed that the material added during that period is dissolved instantaneously.

**Tip:** By setting  $do\_scale=True$ , the composition will be scaled to 100% / fraction of 1. This is useful if the composition provided is not scaling to 100% / 1. An example could be a slag addition which is provided like this: 90 wt-% CaO - 5 wt-% Al2O3 - 4 wt-% SiO2.

# Parameters

- **composition** The composition
- **rate** The rate of addition [kg/s]
- temperature The initial addition temperature (default: 20 °C) [K]
- **composition\_unit** The composition unit
- **do\_scale** If the composition is scaled to 100% / fraction of 1

# $\texttt{get\_composition\_unit()} \rightarrow \textit{CompositionUnit}$

Returns the composition unit used in this addition.

# Returns

The composition unit

# get\_rate() $\rightarrow$ float

Returns the rate of addition.

# Returns

The addition rate [kg/s]

class tc\_python.process\_metallurgy.process.ContinuousGasAddition(composition: Dict[str, float],

rate: float, temperature: float =
293.15, rate\_unit: GasRateUnit
= GasRateUnit.NORM\_CUBIC\_METER\_PER\_SEC,
composition\_unit:
GasCompositionUnit =
GasCompositionUnit.VOLUME\_PERCENT,
do\_scale: bool = False)

# Bases: AbstractContinuousAddition

A gas addition in a process simulation that is added continuously during a period of time.

It is assumed that the gas added during that period is dissolved instantaneously.

**Tip:** By setting  $do\_scale=True$ , the composition will be scaled to 100% / fraction of 1. This is useful if the composition provided is not scaling to 100% / 1. An example could be a gas addition which is provided like this: 90 vol-% Ar - 10 vol-% O2.

# $\texttt{get\_composition\_unit()} \rightarrow \textit{GasCompositionUnit}$

Returns the composition unit used in this addition.

# Returns

The composition unit

## get\_rate() $\rightarrow$ float

Returns the rate of addition.

**Note:** The rate unit can be obtained using *get\_rate\_unit(*).

# Returns

The addition rate [in the rate unit]

 $get_rate_unit() \rightarrow GasRateUnit$ 

Returns the rate unit used in this addition.

#### Returns

The rate unit

class tc\_python.process\_metallurgy.process.ExhaustGasResult(result)

#### Bases: object

A result representing the exhaust gas zone, here all exhaust gas generated during the process is accumulated.

The data is returned for each time point of the process simulation. These time points can be obtained from this method: *ProcessSimulationResult.get\_time\_points()*.

#### $get_amount() \rightarrow List[float]$

Returns the amount of exhaust gas present at each time point.

This is the amount of gas accumulated since the beginning of the process.

#### Returns

The accumulated amount of gas at each time point [kg]

#### 

Returns the amount of each exhaust gas component present at each time point.

This is the amount of gas accumulated since the beginning of the process. This is different from the **current composition at each time point** obtained using *get\_composition()*.

#### Returns

The accumulated amount of each gas component at each time point [kg]

Returns the current composition of the exhaust gas zone at each time point. This is the composition **at each time point**. This is different from the **accumulated amount** obtained using get\_amount\_of\_components().

#### **Parameters**

- **composition\_type** The type of the composition, can be by gas component or by element, **default: by gas component**
- unit The composition unit, default: mass percent

#### Returns

The current composition of the gas components at each time point

#### get\_pressure() $\rightarrow$ List[float]

Returns the pressure of the exhaust gas zone at each time point.

Returns The pressure [Pa]

# $get_stable_phases() \rightarrow Set[str]$

Returns the stable phases within the exhaust gas zone at each time point.

#### Returns

The stable phases

# $\texttt{get\_temperature()} \rightarrow List[float]$

Returns the temperature of the exhaust gas at each time point.

#### Returns

The temperature at each time point [K]

class tc\_python.process\_metallurgy.process.MassTransferCoefficients

# Bases: object

The mass transfer coefficients between a reaction zone and a bulk zone vs. time.

**add**(*mass\_transfer\_coefficient: float, time: float = 0.0*)

Adds the mass transfer coefficient valid beginning at a time point.

This value is valid until another value is defined for a later time point.

# Parameters

- mass\_transfer\_coefficient The mass transfer coefficient [m/s]
- time The time-point where the mass transfer coefficient begins to be valid [s]

#### Returns

This MassTransferCoefficients object

#### class tc\_python.process\_metallurgy.process.MetalBulkZone(density: float)

Bases: Zone

A metallic bulk zone in a process simulation.

This is representing a large volume in the process, for example the steel melt. A zone is a volume in a process that has identical temperature and composition. It has well-defined boundaries to other zones.

The name of this zone is automatically defined and unique.

**add\_addition**(*addition*: AbstractSingleTimeAddition, *time: float* = 0.0)

Adds a single-time addition at the specified time point to the zone. The addition will be dissolved immediately.

# **Parameters**

- addition A SingleTimeAddition or SingleTimeGasAddition
- time The time point [s]

## Returns

This MetalBulkZone object

Adds a constant addition continuously during the specified time period to the zone. All added material will be dissolved immediately.

#### **Parameters**

- addition A ContinuousAddition or ContinuousGasAddition
- from\_time The start time point [s]
- **to\_time** The end time point [s]

# Returns

This MetalBulkZone object

**add\_power**(*power*: *float*, *from\_time*: *float* = 0.0, *to\_time*: *float* = *nan*)

Adds a constant power during a specified time period to the zone (for example heating or cooling).

#### Parameters

- power The power [W]
- from\_time The start time point [s]
- to\_time The end time point [s]

#### Returns

This MetalBulkZone object

# disable\_degassing()

Disables degassing for this zone, i.e. all gas formed at any time step will be staying in this zone.

## Returns

This MetalBulkZone object

# enable\_degassing()

Enables degassing for this zone, i.e. any gas formed at any time step will be removed after that time step. This gas will be transferred into the *exhaust gas zone*. **This is the default**.

#### Returns

This MetalBulkZone object

# $\texttt{get\_density()} \to \texttt{float}$

Returns the density of the zone

## Returns

The density [kg/m\*\*3]

# $\texttt{get\_elements()} \rightarrow Set[str]$

Returns the elements present in the zone. The elements are determined by the additions.

## Returns

The elements

# $\texttt{get\_id}() \to str$

Returns the unique name / id of the zone.

# Returns

The zone name / id

# get\_phase\_group\_to\_transfer() $\rightarrow$ *PhaseGroup*

Returns the phase group that is transferred from the attached reaction zones back to this zone after each time step.

#### Returns

The phase group

#### is\_degassing\_enabled() $\rightarrow$ bool

Returns if degassing is enabled in the zone.

# Returns

If degassing is enabled

```
class tc_python.process_metallurgy.process.ProcessSimulationCalculation(calculation)
```

Bases: object

A Process Metallurgy process simulation. Such calculations represent complete metallurgical processes with several zones and simulate their evolution over time.

 $calculate(timeout_in_minutes: float = 0.0) \rightarrow ProcessSimulationResult$ 

Runs the Process Metallurgy process simulation.

## Parameters

timeout\_in\_minutes - The calculation will be aborted after that time, default: no timeout

# Returns

A new ProcessSimulationResult object

#### set\_end\_time(end\_time: float)

Sets the end time of a process.

## **Parameters** end\_time – The end time point [s]

#### Returns

This ProcessSimulationCalculation object

## set\_initial\_time\_step(initial\_time\_step: float = 1.0)

Sets the initial time step used in the process simulation.

**Note:** All later time steps are automatically determined to limit the expected temperature change during that step, this is controlled by *set\_max\_allowed\_temp\_change\_per\_step()*.

# **Parameters** initial\_time\_step – The initial time step [s]

Returns

This ProcessSimulationCalculation object

# set\_max\_allowed\_temp\_change\_per\_step(max\_allowed\_temp\_change: float = 10.0)

The maximum allowed temperature change per time step. This is implicitly also limiting the composition change during a time step and required for numerical stability.

#### Parameters

max\_allowed\_temp\_change – The maximum allowed temperature change [K]

#### Returns

This ProcessSimulationCalculation object

# set\_max\_time\_step(max\_time\_step: float = 180.0)

The maximum time step chosen by the automatic time step control.

**Note:** All time steps are automatically determined to limit the expected temperature change during that step, this is controlled by *set\_max\_allowed\_temp\_change\_per\_step()*.

#### Parameters

**max\_time\_step** – The maximum time step [s]

## Returns

This ProcessSimulationCalculation object

#### set\_min\_time\_step(min\_time\_step: float = 1.0)

The minimum time step chosen by the automatic time step control.

**Note:** All time steps are automatically determined to limit the expected temperature change during that step, this is controlled by *set\_max\_allowed\_temp\_change\_per\_step()*.

#### **Parameters**

min\_time\_step - The minimum time step [s]

#### Returns

This ProcessSimulationCalculation object

## set\_pressure(pressure: float = 100000.0)

Sets a constant pressure during the complete process.

Parameters pressure – The pressure [Pa]

#### Returns

This ProcessSimulationCalculation object

set\_pressure\_in\_time\_period(pressure\_in\_pa: float, from\_time: float = 0.0, to\_time: float = nan)

Sets a constant pressure during a time period.

# Default: 1.0e5 Pa.

**Parameters** 

- pressure\_in\_pa The pressure [Pa]
- from\_time The start time [s]
- **to\_time** The end time [s]

#### Returns

This *ProcessSimulationCalculation* object

#### with\_options(options: ProcessMetallurgyOptions)

Sets the options for the process simulation.

# Parameters

**options** – The options

# Returns

This ProcessSimulationCalculation object

## with\_reaction\_zone(reaction\_zone: ReactionZone)

Sets the reaction zone of the process simulation. The bulk zones attached to this reaction zone are configured in the reaction zone object.

Note: In the present release, only one reaction zone is supported.

#### **Parameters**

reaction\_zone – The reaction zone object

#### Returns

This ProcessSimulationCalculation object

#### class tc\_python.process\_metallurgy.process.ProcessSimulationResult(result)

Bases: AbstractResult

The result of a Process Metallurgy process simulation.

# $get\_activity\_of\_slag(zone: Union[Zone, str], component: str, reference: ActivityReference = ActivityReference.LIQUID) \rightarrow List[float]$

Returns the activity of a component in the slag in a zone at each time point.

## **Parameters**

- **zone** The zone object or the zone name
- component The component
- **reference** The reference for the activity, can be liquid or solid slag, **default: liquid slag**

## Returns

The activity of the component at each time point [-]

## get\_amount(*zone: Union*[Zone, *str*]) → List[float]

Returns the amount of a zone at each time point.

#### **Parameters**

**zone** – The zone object or the zone name

#### Returns

The amount at each time point [kg]

# $\texttt{get\_amount\_of\_elements()} \rightarrow Dict[str, List[float]]$

Returns the total amount of each element in the simulation at each time point.

#### Returns

The total amount of the elements at each time point [kg]

# 

Returns the amount of each phase group (e.g., for example all liquid slag) in a zone at each time point.

# Parameters

**zone** – The zone object or the zone name

# Returns

The amount of the phase groups at each time point [kg]

# get\_amount\_of\_phases(zone: Union[Zone, str]) → Dict[str, List[float]]

Returns the amount of each phase in a zone at each time point.

#### Parameters

zone - The zone object or the zone name

#### Returns

The amount of the phases at each time point [kg]

# $get\_components() \rightarrow Set[str]$

Returns all components defined in the simulation.

# Returns

The components

get\_composition(zone: Union[Zone, str], composition\_unit: CompositionUnit = CompositionUnit.MASS\_PERCENT) → Dict[str, List[float]]

Returns the composition of a zone per element at each time point.

# Parameters

- **zone** The zone object or the zone name
- composition\_unit The composition unit, default: mass percent

# Returns

The composition at each time point

get\_composition\_of\_phase(zone: Union[Zone, str], phase: str, composition\_unit: CompositionUnit = CompositionUnit.MASS\_PERCENT, composition\_type: CompositionType = CompositionType.COMPONENT) → Dict[str, List[float]]

Returns the composition of a phase in a zone at each time point.

# Parameters

- zone The zone object or the zone name
- phase The phase name
- composition\_unit The composition unit, default: mass percent
- **composition\_type** Defines if the composition is given by element (e.g., 75 *wt-% Fe* 25 *wt-% Cr*) or by component (e.g. 65 *wt-% Al2O3 35 wt-% CaO*). In case of a metallic phase, the composition is given by element even if *component* is selected. **Default: by component**.

# Returns

The composition at each time point

CompositionType = CompositionType.COMPONENT)  $\rightarrow$  Dict[str, List[float]]

Returns the composition of a phase group (e.g., all liquid slag) in a zone at each time point.

# Parameters

- zone The zone object or the zone name
- **phase\_group** The phase group
- composition\_unit The composition unit, default: mass percent
- **composition\_type** Defines if the composition is given by element (e.g., 75 wt-% Fe 25 wt-% Cr) or by component (e.g. 65 wt-% Al2O3 35 wt-% CaO). In case of a metallic phase, the composition is given by element even if *component* is selected. **Default: by component**.

# Returns

The composition at each time point

#### $get_elements() \rightarrow Set[str]$

Returns all elements present in the simulation.

# Returns

The elements

# $\texttt{get\_enthalpy()} \rightarrow List[float]$

Returns the total enthalpy of the process at each time point.

#### Returns

The enthalpy at each time point [J]

# $get_exhaust_gas() \rightarrow ExhaustGasResult$

Returns the result for the exhaust gas zone.

This result object can be used to evaluate the exhaust gas zone at each time point.

#### Returns

The exhaust gas zone result object.

Returns the Thermo-Calc Console syntax formula used for calculating the activity of a component in the slag (e.g. *AC(AL2O3, IONIC\_LIQ)* in a zone at each time point. The actual activity can be obtained using *get\_activity\_of\_slag()*.

## Parameters

- **zone** The zone object or the zone name
- **component** The component
- **reference** The reference for the activity, can be liquid or solid slag, **default: liquid slag**

### Returns

The formula for calculating the activity at each time point

 $\label{eq:get_formula_for_slag_property} (zone: Union[Zone, str], slag_property: SlagProperty, slag_type: \\ SlagType = SlagType.ALL) \rightarrow List[str]$ 

Returns the Thermo-Calc Console syntax formula used for calculating a property of the slag (e.g. B(CAO)/B(SIO2) in a zone at each time point. The actual slag property can be obtained using  $get\_slag\_property()$ .

#### Parameters

- zone The zone object or the zone name
- slag\_property The slag property
- **slag\_type** The part of the slag for which the property will be calculated. Can be all slag, the liquid or the solid slag. **Default: all slag**

#### Returns

The formula for calculating the slag property at each time point

```
get_fraction_of_phase_groups(zone: Union[Zone, str], unit: PhaseUnit = PhaseUnit.MASS_FRACTION) \rightarrow Dict[PhaseGroup, List[float]]
```

Returns the fractions of the phase groups (e.g., all liquid slag) in a zone at each time point.

#### **Parameters**

• **zone** – The zone object or the zone name

• unit – The unit of the fraction

#### Returns

The phase fractions at each time point

```
get_fraction_of_phases(zone: Union[Zone, str], unit: PhaseUnit = PhaseUnit.MASS_FRACTION) \rightarrow Dict[str, List[float]]
```

Returns the fractions of all stable phases in a zone at each time point.

#### Parameters

- **zone** The zone object or the zone name
- unit The unit of the fraction

#### Returns

The phase fractions at each time point

#### $\texttt{get\_gas\_components()} \rightarrow \texttt{Set[str]}$

Returns all components of the gas phase defined for the elements present in the simulation.

#### Returns

The components of the gas phase

# $\texttt{get\_num\_of\_performed\_steps()} \rightarrow List[int]$

Returns the accumulated number of performed time steps at each time point.

**Note:** The number of performed time steps can differ from the index of the time step in the result list because time steps might have been repeated with smaller step size during a process simulation.

# Returns

The accumulated number of performed time steps

#### 

Returns the partial pressure of oxygen in the zone at each time point.

# Parameters

**zone** – The zone object or the zone name

# Returns

The partial pressure [Pa]

# get\_pressure(zone: Union[Zone, str]) $\rightarrow$ List[float]

Returns the pressure in a zone at each time point.

# Parameters

**zone** – The zone object or the zone name

# Returns

The pressure at each time point [Pa]

**get\_slag\_property**(*zone: Union*[Zone, *str*], *slag\_property:* SlagProperty, *slag\_type:* SlagType = SlagType.ALL) → List[float]

Returns a property of the slag in a zone at each time point. These properties are mostly used to describe the property of a slag to pick up sulfur.

# Parameters

• zone – The zone object or the zone name

- slag\_property The slag property
- **slag\_type** The part of the slag for which the property will be calculated. Can be all slag, the liquid or the solid slag. **Default: all slag**

## Returns

The slag property at each time point [unit depending on the property]

get\_stable\_phases(zone: Union[Zone, str]) → Set[str]

Returns the stable phases in a zone.

#### Parameters

zone - The zone object or the zone name

## Returns

The stable phases

 $\texttt{get\_stable\_phases\_in\_phase\_group}(\textit{zone: Union[Zone, str]}, \textit{phase\_group: PhaseGroup}) \rightarrow \texttt{Set[str]}$ 

Returns the stable phases of a phase group (e.g., all solid slag) in a zone.

#### Parameters

- zone The zone object or the zone name
- phase\_group The phase group

#### Returns

The stable phases of the phase group

# get\_temperature(*zone: Union*[Zone, *str*]) $\rightarrow$ List[float]

Returns the temperature of a zone at each time point.

#### Parameters

**zone** – The zone object or the zone name

# Returns

The temperature at each time point [K]

#### get\_time\_points() → List[float]

Returns the time points of the process simulation. All result quantities are returned for exactly these time points.

#### Returns

The time points [s]

**get\_value\_of**(*zone: Union*[Zone, *str*], *classic\_expression: str*) → List[float]

Returns a value for a thermodynamic quantity in a zone at each time point.

**Warning:** It should normally not be required to use this method, use the appropriate method available in the API instead.

#### **Parameters**

- zone The zone object or the zone name
- **classic\_expression** The thermodynamic quantity to get the value of in Thermo-Calc Console Mode syntax (for example "NPM(FCC\_A1)")

#### Returns

The requested value at each time point

```
get_viscosity_dynamic_of_phase(zone: Union[Zone, str], phase: str) → List[float]
```

Returns the dynamic viscosity of a phase in a zone at each time point.

## **Parameters**

- zone The zone object or the zone name
- **phase** The phase name

Returns

The dynamic viscosity at each time point [Pa\*s]

#### get\_viscosity\_kinematic\_of\_phase(zone: Union[Zone, str], phase: str) → List[float]

Returns the kinematic viscosity of a phase in a zone at each time point.

#### Parameters

- zone The zone object or the zone name
- **phase** The phase name

## Returns

The kinematic viscosity at each time point [m\*\*2/s]

class tc\_python.process\_metallurgy.process.ReactionZone(area: float, left\_zone: Zone,

mass\_transfer\_coefficient\_left: Union[float, MassTransferCoefficients], right\_zone: Zone, mass\_transfer\_coefficient\_right: Union[float, MassTransferCoefficients])

## Bases: Zone

A reaction zone in a process simulation, this is representing the interface layer between two bulk zones that are in contact and can react with each other, for example the steel melt and the top slag. The size of the reaction zone is dynamic and determined by the mass transfer coefficient. A zone is a volume in a process that has identical temperature and composition. It has well-defined boundaries to other zones.

add\_addition(addition: AbstractSingleTimeAddition, time: float = 0.0)

Adds a single-time addition at the specified time point to the zone. The addition will be dissolved immediately.

# **Parameters**

- addition A SingleTimeAddition or SingleTimeGasAddition
- time The time point [s]

# Returns

This *ReactionZone* object

Adds a constant addition continuously during the specified time period to the zone. All added material will be dissolved immediately.

#### Parameters

- addition A ContinuousAddition or ContinuousGasAddition
- from\_time The start time point [s]
- **to\_time** The end time point [s]

#### Returns

This ReactionZone object

# add\_heat\_transfer(heat\_transfer\_coefficient: float)

Adds heat transfer through the reaction zone, i.e., between the two attached bulk zones.

# Parameters

heat\_transfer\_coefficient – The heat transfer coefficient [W/(K\*m\*\*2)]

#### Returns

This ReactionZone object

**add\_power**(*power: float, from\_time: float = 0.0, to\_time: float = nan*)

Adds a constant power during a specified time period to the zone (for example heating or cooling).

# Parameters

- **power** The power [W]
- **from\_time** The start time point [s]
- to\_time The end time point [s]

#### Returns

This ReactionZone object

# add\_transfer\_of\_phase\_group(transfer\_of\_phase\_group: TransferOfPhaseGroup)

Adds transfer of a certain phase group through the reaction zone during each time step, i.e. from one of the attached bulk zones to the other. This is for example used to model inclusion flotation from the steel melt to the slag.

#### **Parameters**

**transfer\_of\_phase\_group** – The transfer of phase group configuration, can be time-dependent.

# Returns

This ReactionZone object

# disable\_degassing()

Disables degassing for this zone, i.e. all gas formed at any time step will be staying in this zone.

# Returns

This ReactionZone object

# enable\_degassing()

Enables degassing for this zone, i.e. any gas formed at any time step will be removed after that time step. This gas will be transferred into the *exhaust gas zone*. **This is the default**.

# Returns

This ReactionZone object

# $\texttt{get\_elements()} \rightarrow Set[str]$

Returns the elements present in the zone. The elements are determined by the additions.

# Returns

The elements

# $\texttt{get\_id}() \to str$

Returns the unique id of the zone. :return: The zone id

# $\texttt{is\_degassing\_enabled()} \rightarrow \texttt{bool}$

Returns if degassing is enabled in the zone.

# Returns

If degassing is enabled

**class** tc\_python.process\_metallurgy.process.**SingleTimeAddition**(*composition: Dict[str, float]*,

amount: float, temperature: float = 293.15, composition\_unit: CompositionUnit = CompositionUnit.MASS\_PERCENT, do\_scale: bool = False)

Bases: AbstractSingleTimeAddition

An addition in a process simulation that is added at a distinct time point.

It is assumed that the addition is dissolved instantaneously.

**Tip:** By setting  $do\_scale=True$ , the composition will be scaled to 100% / fraction of 1. This is useful if the composition provided is not summing to 100% / 1. An example could be a slag addition which is provided like this: 90 wt-% CaO - 5 wt-% Al2O3 - 4 wt-% SiO2.

# Parameters

- composition The composition
- amount The amount [kg]
- temperature The initial addition temperature (default: 20 °C) [K]
- composition\_unit The composition unit
- **do\_scale** If the composition is scaled to 100% / fraction of 1

## $\texttt{get\_amount()} \to \texttt{float}$

Returns the amount of this addition.

## Returns

The amount [kg]

## $get\_composition\_unit() \rightarrow CompositionUnit$

Returns the composition unit used in this addition.

# Returns

The composition unit

class tc\_python.process\_metallurgy.process.SingleTimeGasAddition(composition: Dict[str, float],

amount: float, temperature: float = 293.15, amount\_unit: GasAmountUnit = GasAmountUnit.NORM\_CUBIC\_METER, composition\_unit: GasCompositionUnit = GasCompositionUnit.VOLUME\_PERCENT, do\_scale: bool = False)

Bases: AbstractSingleTimeAddition

A gas addition in a process simulation that is added at a distinct time point.

It is assumed that the addition is dissolved instantaneously.

**Tip:** By setting  $do\_scale=True$ , the composition will be scaled to 100% / fraction of 1. This is useful if the composition provided is not scaling to 100% / 1. An example could be a gas addition which is provided like this: 90 vol-% Ar - 10 vol-% O2.

#### Parameters

- **composition** The composition
- **amount** The amount
- temperature The initial addition temperature (default: 20 °C) [K]
- **amount\_unit** The amount unit
- **composition\_unit** The composition unit
- do\_scale If the composition is scaled to 100% / fraction of 1

# $\texttt{get\_amount()} \rightarrow \texttt{float}$

Returns the amount of this addition.

Note: The amount unit can be obtained using get\_amount\_unit().

#### Returns

The amount [in the amount unit]

#### $get\_amount\_unit() \rightarrow GasAmountUnit$

Returns the amount unit used in this addition.

Returns

The amount unit

# $get\_composition\_unit() \rightarrow GasCompositionUnit$

Returns the composition unit used in this addition.

# Returns

The composition unit

#### class tc\_python.process\_metallurgy.process.SlagBulkZone(density: float)

Bases: Zone

A slag bulk zone in a process simulation.

This is representing a large volume in the process, for example the top slag. A zone is a volume in a process that has identical temperature and composition. It has well-defined boundaries to other zones.

The name of this zone is automatically defined and unique.

**add\_addition**(*addition*: AbstractSingleTimeAddition, *time: float = 0.0*)

Adds a single-time addition at the specified time point to the zone. The addition will be dissolved immediately.

#### **Parameters**

- addition A SingleTimeAddition or SingleTimeGasAddition
- time The time point [s]
#### Returns

This SlagBulkZone object

Adds a constant addition continuously during the specified time period to the zone. All added material will be dissolved immediately.

#### **Parameters**

- addition A ContinuousAddition or ContinuousGasAddition
- **from\_time** The start time point [s]
- **to\_time** The end time point [s]

### Returns

This SlagBulkZone object

#### **add\_power**(*power*: *float*, *from\_time*: *float* = 0.0, *to\_time*: *float* = *nan*)

Adds a constant power during a specified time period to the zone (for example heating or cooling).

#### **Parameters**

- power The power [W]
- **from\_time** The start time point [s]
- to\_time The end time point [s]

#### Returns

This SlagBulkZone object

# disable\_degassing()

Disables degassing for this zone, i.e. all gas formed at any time step will be staying in this zone.

#### Returns

This SlagBulkZone object

# enable\_degassing()

Enables degassing for this zone, i.e. any gas formed at any time step will be removed after that time step. This gas will be transferred into the *exhaust gas zone*. **This is the default**.

# Returns

This SlagBulkZone object

# $\texttt{get\_density()} \to \texttt{float}$

Returns the density of the zone

# Returns

The density [kg/m\*\*3]

#### $\texttt{get\_elements()} \rightarrow Set[str]$

Returns the elements present in the zone. The elements are determined by the additions.

#### Returns

The elements

# $get_id() \rightarrow str$

Returns the unique id of the zone. :return: The zone id

### get\_phase\_group\_to\_transfer() $\rightarrow$ *PhaseGroup*

Returns the phase group that is transferred from the attached reaction zones back to this zone after each time step.

#### Returns

The phase group

# $\texttt{is\_degassing\_enabled()} \rightarrow \texttt{bool}$

Returns if degassing is enabled in the zone.

# Returns

If degassing is enabled

### Bases: object

The transfer of a percentage of a certain phase group (.e.g., solid slag) between zones during each time step. This is for example used to model inclusion flotation from the steel melt to the slag.

# **add**(*transfer\_rate: float*, *time: float* = 0)

Adds the transfer rate valid beginning at a time point.

This value is valid until another value is defined for a later time point.

### **Parameters**

- transfer\_rate The transfer rate [% of phase group amount/s]
- time The time point where the transfer of a phase group begins to be valid [s]

### Returns

This TransferOfPhaseGroup object

### get\_phase\_group\_to\_transfer() $\rightarrow$ *PhaseGroup*

Returns the phase group to be transferred

### Returns

The phase group

### get\_transfer\_source\_zone\_id() $\rightarrow$ str

The id of the source zone of the transfer

### Returns

This source zone id

### class tc\_python.process\_metallurgy.process.Zone

Bases: object

The base class of a zone in a process simulation. A zone is a volume in a process that has identical temperature and composition. It has well-defined boundaries to other zones.

add\_addition(addition: AbstractSingleTimeAddition, time: float = 0.0)

Adds a single-time addition at the specified time point to the zone. The addition will be dissolved immediately.

### **Parameters**

- addition A SingleTimeAddition or SingleTimeGasAddition
- **time** The time point [s]

#### Returns

This Zone object

Adds a constant addition continuously during the specified time period to the zone. All added material will be dissolved immediately.

#### **Parameters**

- addition A ContinuousAddition or ContinuousGasAddition
- **from\_time** The start time point [s]
- **to\_time** The end time point [s]

### Returns

This Zone object

#### **add\_power**(*power*: *float*, *from\_time*: *float* = 0.0, *to\_time*: *float* = *nan*)

Adds a constant power during a specified time period to the zone (for example heating or cooling).

#### **Parameters**

- power The power [W]
- **from\_time** The start time point [s]
- to\_time The end time point [s]

#### Returns

This Zone object

# disable\_degassing()

Disables degassing for this zone, i.e. all gas formed at any time step will be staying in this zone.

#### Returns

This Zone object

# enable\_degassing()

Enables degassing for this zone, i.e. any gas formed at any time step will be removed after that time step. This gas will be transferred into the *exhaust gas zone*. **This is the default**.

# Returns

This Zone object

#### $get_elements() \rightarrow Set[str]$

Returns the elements present in the zone. The elements are determined by the additions.

### Returns

The elements

# $\texttt{get\_id}() \to str$

Returns the unique id of the zone. :return: The zone id

#### is\_degassing\_enabled() $\rightarrow$ bool

Returns if degassing is enabled in the zone.

#### Returns

If degassing is enabled

# 6.1.10 Module "additive\_manufacturing"

# class tc\_python.am.AdditiveManufacturingCalculation(calculation)

Bases: object

Abstract base class for an Additive Manufacturing calculation.

# disable\_fluid\_flow\_marangoni()

Disables the fluid flow modelling of the Marangoni effect.

Default: Enabled

# Returns

This AdditiveManufacturingCalculation object

# disable\_separate\_materials()

Disables separate material properties for powder and solid material.

# Default: Disabled

# Returns

This AdditiveManufacturingCalculation object

# enable\_fluid\_flow\_marangoni()

Enables the fluid flow modelling of the Marangoni effect.

# Default: Enabled

**Note:** This option is not possible to use in conjunction with the option *separate material*, which is therefore automatically disabled.

# Returns

This AdditiveManufacturingCalculation object

# enable\_separate\_materials()

Enables separate material properties for powder and solid material.

# Default: Disabled

**Note:** This option is not possible to use in conjunction with the option *Marangoni fluid flow*, which is therefore automatically disabled.

# Returns

This AdditiveManufacturingCalculation object

# $\texttt{get\_configuration\_as\_string()} \to \operatorname{str}$

Returns detailed information about the current state of the calculation object.

**Warning:** The structure of the calculation objects is an implementation detail and might change between releases without notice. **Therefore do not rely on the internal object structure**.

# invalidate()

Invalidates the object and frees the disk space used by it. *This is only required if the disk space occupied by the object needs to be released during the calculation.* No data can be retrieved from the object afterward.

set\_ambient\_temperature(temperature: float = 296.15)

Sets the ambient temperature.

Default: 23 degree Celsius

#### Parameters

temperature – The ambient temperature [K]

#### Returns

This AdditiveManufacturingCalculation object

#### set\_base\_plate\_temperature(temperature: float = 303.15)

Sets the baseplate temperature.

Default: 30 degree Celsius

# Parameters

temperature – The baseplate temperature [K]

#### Returns

This AdditiveManufacturingCalculation object

#### set\_gas\_pressure(pressure: float = 100000.0)

Sets the gas pressure.

Default: 1.0e5 Pa

# Parameters pressure – The pressure [Pa]

Returns This AdditiveManufacturingCalculation object

# set\_height(height: float = 0.002)

Sets the height of the simulation domain.

#### Default: 2.0e-3 m

Parameters height – The height [m]

### Returns

This AdditiveManufacturingCalculation object

# set\_layer\_thickness(layer\_thickness: float = 4e-05)

Sets the layer thickness.

Default: 40.0e-6 m

#### Parameters

layer\_thickness – The layer thickness [m]

#### Returns

This AdditiveManufacturingCalculation object

#### set\_powder\_density(powder\_density: float = 80.0)

Sets the powder density.

**Default:** 80.0%

Parameters powder\_density – The powder density [%]

#### Returns

This AdditiveManufacturingCalculation object

# with\_heat\_source(heat\_source: HeatSource)

Sets the heat source.

Parameters heat\_source – The heat source

#### Returns

This AdditiveManufacturingCalculation object

### with\_material\_properties(material\_properties: MaterialProperties)

Sets the material properties.

**Tip:** Material properties can be defined like this: MaterialProperties.from\_library("IN718") or MaterialProperties.from\_scheil\_result(scheil\_result).

#### Parameters

material\_properties - The material properties

#### Returns

This AdditiveManufacturingCalculation object

### with\_mesh(mesh: Mesh)

Sets the mesh.

### Parameters mesh – The mesh

#### Returns

This AdditiveManufacturingCalculation object

### with\_numerical\_options(numerical\_options: NumericalOptions)

Sets the numerical options.

# Parameters

numerical\_options – The numerical options

### Returns

This AdditiveManufacturingCalculation object

### with\_top\_boundary\_conditions(boundary\_conditions: TopBoundaryConditions)

Sets the boundary conditions.

### Parameters

boundary\_conditions – The boundary conditions

### Returns

This AdditiveManufacturingCalculation object

### class tc\_python.am.AdditiveManufacturingResult(result)

Bases: AbstractResult

Base class for additive manufacturing results.

#### get\_absorptivity\_mesh()

Returns a mesh object containing the absorptivity data from this result. Returns NONE for this method if the result has no absorptivity data.

**get\_data\_over\_line**(*name: str, point1: List[float], point2: List[float], n: int*) → Tuple[List[float], List[float]]

Obtains the data of property name over a line from point1 to point2 :param name: The name of property to obtain the data from :param point1: The first point of the line :param point2: The second point of the line :param n: The number of points to obtain :return: A tuple (distance to point 1 [m], data value)

#### get\_effective\_absorption() $\rightarrow$ float

:return Effective absorption under the heat source due to multiple reflections of laser rays

Returns a pyvista mesh object that can be added to a pyvista plotter.

More details about *pyvista* meshes can be found in their documentation: https://docs.pyvista.org/version/ stable/api/plotting/\_autosummary/pyvista.Plotter.add\_mesh.html

**Tip:** This method is typically used to obtain additional meshes with other settings if the plot object has already been retrieved with *get\_pyvista\_plotter()*.

#### **Parameters**

- scalar The quantity to be visualized in the plot
- **material\_type** The material type to be visualized in the plot

#### Returns

A pyvista.DataSet object

```
get_pyvista_plotter(scalar: Scalar = Scalar.TEMPERATURE, material_type: MaterialType =
    MaterialType.SHOW_ALL, camera: Optional[Dict[str, List[float]]] = None,
    anti_aliasing: str = 'msaa', multi_samples: int = 16, color_map: str = 'jet',
    enable_camera_orientation_widget: bool = False, render_lines_as_tubes: bool =
    True, background: str = 'LightSteelBlue', shape: Optional[bool] = None,
```

view\_buttons: bool = True, update\_plot\_callback: Optional[Callable] = None)

Returns a *pyvista* plotter and a mesh object containing the data from this result. They can be used to create 3D-plots visualizing the results of the AM calculation. More details about the *pyvista* settings can be found in their documentation: https://docs.pyvista.org/version/stable/api/plotting/\_autosummary/pyvista.Plotter. html

The most simple usage of this method is:

```
plotter, mesh = result.get_pyvista_plotter()
plotter.add_mesh(mesh)
plotter.show()
```

#### Parameters

- scalar The quantity to be visualized in the plot
- material\_type The material type to be visualized in the plot, only available if AdditiveManufacturingCalculation.enable\_separate\_materials() has been used

- **camera** Defining the camera position, view-up vector and focus point, the default is DEFAULT\_CAMERA, for example: { 'position': [-2, -2, 1], 'viewup': [0, 0, 1], 'focal\_point': [0.0, 0.0, 0.0]}. More details can be found here: https://docs.pyvista.org/version/stable/ api/core/camera.html
- anti\_aliasing The *pyvista* antialiasing setting, can be one of: "*msaa*" Multi-Sample Anti-Aliasing, "*fxaa*" Fast Approximate Anti-Aliasing or "*ssaa*" Super-Sample Anti-Aliasing. More details can be found here: https://docs.pyvista.org/version/stable/examples/02-plot/anti-aliasing.html
- multi\_samples The number of samples used for antialiasing
- color\_map The *pyvista* colormap. Can be any colormap provided by Matplotlib and some other plotting libraries. More details can be found here: https://docs.pyvista.org/ version/stable/examples/02-plot/cmap.html
- **enable\_camera\_orientation\_widget** Enables the *pyvista* camera orientation widget in the plotter
- render\_lines\_as\_tubes Controls if lines are rendered as tubes
- **background** The *pyvista* background color, either a string, rgb list or hex color string. https://docs.pyvista.org/version/stable/api/plotting/\_autosummary/pyvista.Plotter. background\_color.html
- **shape** The shape of the plot, i.e. how many subplots will be created a tuple (y, x), for example (2, 2). They are accessed using pyvista.Plotter.subplot.
- **view\_buttons** If buttons for quick navigation between camera and x-, y-, or z-direction view should be added to the plot
- update\_plot\_callback A plot update function that will be called every time the slider with simulation time is dragged in the plot windows, this can be used to dynamically apply changes to the plot, required syntax: def update\_plot(plotter: pv.Plotter, mesh: pv.DataSet)

#### Returns

Tuple containing the pyvista.Plotter and the pyvista.DataSet mesh object

### get\_result\_file\_path() → str

Returns the path to the main result file (for example a ParaView \*.pvd file) on disk containing the complete result data set. Its directory contains also further result data that can be useful.

**Tip:** The Python API of ParaView can be used to extract any kind of data from the result in a programmatic way. See here for more details: https://kitware.github.io/paraview-docs/latest/python/

#### Returns

The path to the main result file

# $\texttt{get\_thermal\_gradient\_and\_melting\_rate()} \rightarrow List[List[float]]$

Evaluate the thermal gradient and melting rate at the liquidus isotherm :return [[Thermal gradient], [melting rate], [x], [y], [z]]

### $\texttt{get\_thermal\_gradient\_and\_solidification\_rate()} \rightarrow \texttt{List[List[float]]}$

Evaluate the thermal gradient and solidification rate at the liquidus isotherm :return [[Thermal gradient], [solidification rate], [x], [y], [z]]

### $\texttt{has\_absorption\_distribution\_on\_keyhole()} \rightarrow \texttt{bool}$

:return True if there is effective absorption on the keyhole

#### class tc\_python.am.Automatic

Bases: FileSavingStrategy

An automatic saving strategy.

#### $\texttt{get\_type()} \to \textit{str}$

Returns the type of the file saving strategy.

Returns

The type

### set\_max\_number\_of\_files\_stored(max\_number: int = 1000000)

Sets the maximum number of files that are stored.

#### **Default: unlimited**

#### Parameters

max\_number – The maximum number of files that is stored

#### Returns

This FileSavingStrategy object

# $\verb|set_saving_interval\_strategy(saving_interval\_strategy=AutomaticSavingIntervalStrategy.LINEAR)|$

Sets the saving interval strategy.

### Default: A linear saving interval strategy.

# Parameters

### saving\_interval\_strategy - The saving interval strategy

#### Returns

This FileSavingStrategy object

# store\_unlimited\_number\_of\_files()

Sets the maximum number of files that are stored to unlimited.

#### Returns

This FileSavingStrategy object

### class tc\_python.am.AutomaticSavingIntervalStrategy(value)

Bases: Enum

The strategy for choosing the time interval for saving files in *automatic* mode.

### **EXPONENTIALLY\_INCREASING = 2**

An exponentially increasing time interval

# LINEAR = 1

A linear time interval

# class tc\_python.am.BiDirectionalScanningStrategy

Bases: ScanningStrategy

A bidirectional scanning strategy (flipping scanning direction of the heat source between alternate tracks).

#### get\_type() → str

Returns the type of scanning strategy.

Returns The type

# set\_angle(angle: float = 0.0)

Sets the rotation of the scanning direction between two consecutive layers.

Note: The scanning direction of the first layer is always aligned parallel to the x-axis.

Default: 0 degree

Parameters angle – The angle [degree]

Returns

This BiDirectionalScanningStrategy object

### set\_hatch\_spacing(hatch\_spacing: float = 0.0)

Sets the horizontal separation between two consecutive tracks.

Default: 0 m

**Parameters** hatch\_spacing – The hatch spacing [m]

Returns

This BiDirectionalScanningStrategy object

# set\_lift\_time(lift\_time: float = 0.0)

Sets the lift time, i.e. the time between two tracks where the heat source is inactive.

Default: 0 s

Parameters lift\_time – The lift time [s]

#### Returns

This BiDirectionalScanningStrategy object

### set\_margin(margin: float = 0.001)

Sets the margin.

This is the offset of the scanning path from the sides of the computational domain.

**Default**: 1.0e-3 [m]

Parameters

margin – The margin [m]

#### Returns

This BiDirectionalScanningStrategy object

set\_number\_of\_layers(number\_of\_layers: int = 1)

Sets the number of layers.

# Default: 1

Parameters number\_of\_layers – The number of layers

# Returns

This BiDirectionalScanningStrategy object

set\_powder\_fill\_time(powder\_fill\_time: float = 0.0)

Sets the powder fill time.

Default: 0 s

Parameters
 powder\_fill\_time - The powder fill time [s]

Returns

This BiDirectionalScanningStrategy object

 $\label{eq:class_$ 

1064.0)

Bases: object

A CoreRing heat source (with calculated absorptivity)

### disable\_keyhole\_model()

Disables using a keyhole model in the simulation.

Returns

This CalculatedCoreRingHeatSource object

static get\_type()  $\rightarrow$  str

Returns the type of heat source.

Returns

The type

set\_absorptivity\_pre\_factor(pre\_factor: float = 1)

**Parameters pre\_factor** – The absorptivity prefactor value to set, default is 1.

Returns

This CalculatedCoreRingHeatSource object

set\_beam\_radius\_core(beam\_radius\_core: float = 5e-05)

Sets the beam radius-Core.

Default: 50.0e-6 m

## Parameters

beam\_radius\_core - The beam radius of the core beam [m]

Returns

This CoreRingHeatSource object

set\_beam\_radius\_ring(beam\_radius\_ring: float = 3.3e-05)

Sets the beam radius-Ring.

Default: 33.0e-6 m

**Note:** set\_beam\_radius\_ring() cannot be used with set\_core\_ring\_index().

#### Parameters

**beam\_radius\_ring** – The beam radius of the ring beam [m]

Returns

This CoreRingHeatSource object

set\_core\_ring\_index(core\_ring\_index: CoreRingIndex = CoreRingIndex.INDEX\_0)

Sets the *Core-Ring Index. Beam radius-Ring, Ring radius* and *Percentage of power in ring* are then set to a value based on the selected Index and Beam radius-Core.

**Default**: CoreRingIndex.INDEX\_0

**Note:** set\_core\_ring\_index() cannot be used with set\_beam\_radius\_ring(),
set\_ring\_radius() or set\_ring\_power\_percent().

Parameters

core\_ring\_index - The index (INDEX\_0 - INDEX\_6) of the Core-Ring laser system

#### Returns

This CoreRingHeatSource object

### set\_power(power: float = 120.0)

Sets the power of the heat source.

Default: 120 W

Parameters power – The power [W]

Returns

This CalculatedCoreRingHeatSource object

set\_ring\_power\_percent(ring\_power\_percent: float = 7.0)

Sets the percentage of the power of the heat source in the ring beam

**Default**: 7.0 %

**Note:** set\_ring\_power\_percent() cannot be used with set\_core\_ring\_index().

Parameters

**ring\_power\_percent** – The percentage of power in the ring bean [%]

#### Returns

This CoreRingHeatSource object

#### set\_ring\_radius(ring\_radius: float = 9.858e-05)

Sets the ring radius.

Default: 98.58e-6 m

**Note:** set\_ring\_radius() cannot be used with set\_core\_ring\_index().

# Parameters

ring\_radius – The radius of the ring beam [m]

# Returns

This CoreRingHeatSource object

set\_scanning\_speed(beam\_speed: float = 0.5)

Sets the moving velocity of the heat source.

Default: 500.0e-3 m/s

Parameters beam\_speed – The beam speed [m/s]

#### Returns

This CalculatedCoreRingHeatSource object

set\_wave\_length(wave\_length: float = 1064)

#### Parameters

wave\_length - The wavelength to be set for the heat source, defaults to 1064 nm.

#### Returns

This CalculatedCoreRingHeatSource objectT

# with\_keyhole\_model(config: KeyholeModel)

Sets the keyhole model applied in the simulation.

# **Default: None**

Parameters config – The keyhole model

#### Returns

This CalculatedCoreRingHeatSource object

**class** tc\_python.am.**CalculatedGaussianHeatSource**(*absorptivity\_factor: float = 1.0, wave\_length: float =* 

1064.0)

Bases: object

A Gaussian heat source (with calculated absorptivity)

# disable\_keyhole\_model()

Disables using a keyhole model in the simulation.

# Returns

This GaussianHeatSource object

# static get\_type() $\rightarrow$ str

Returns the type of heat source.

#### Returns

The type

### set\_absorptivity\_pre\_factor(pre\_factor: float = 1)

### Parameters

**pre\_factor** – The absorptivity prefactor value to set, default is 1.

# Returns

This GaussianHeatSource object

### set\_beam\_radius(beam\_radius: float = 0.00011)

Sets the beam radius.

### **Default**: 110.0e-6 m

# Parameters beam\_radius – The beam radius [m]

#### Returns

This GaussianHeatSource object

### set\_power(power: float = 120.0)

Sets the power of the heat source.

Default: 120 W

### Parameters

**power** – The power [W]

#### Returns

This GaussianHeatSource object

### set\_scanning\_speed(beam\_speed: float = 0.5)

Sets the moving velocity of the heat source.

# Default: 500.0e-3 m/s

Parameters beam\_speed – The beam speed [m/s]

## Returns

This GaussianHeatSource object

set\_wave\_length(wave\_length: float = 1064)

#### Parameters

wave\_length – The wavelength to be set for the heat source, defaults to 1064 nm.

#### Returns

This GaussianHeatSource objectT

### with\_keyhole\_model(config: KeyholeModel)

Sets the keyhole model applied in the simulation.

Default: None

Parameters config – The keyhole model

#### Returns

This GaussianHeatSource object

**class** tc\_python.am.**CalculatedTophatHeatSource**(*absorptivity\_factor: float = 1.0, wave\_length: float =* 

1064.0)

Bases: object

A Tophat heat source (with calculated absorptivity)

# disable\_keyhole\_model()

Disables using a keyhole model in the simulation.

#### Returns

This CalculatedTophatHeatSource object

#### static get\_type() $\rightarrow$ str

Returns the type of heat source.

#### Returns

The type

set\_absorptivity\_pre\_factor(pre\_factor: float = 1)

## Parameters

**pre\_factor** – The absorptivity prefactor value to set, default is 1.

#### Returns

This CalculatedTophatHeatSource object

set\_beam\_radius(beam\_radius: float = 0.00011)

Sets the beam radius.

Default: 110.0e-6 m

Parameters beam\_radius – The beam radius [m]

### Returns

This CalculatedTophatHeatSource object

# set\_power(power: float = 120.0)

Sets the power of the heat source.

Default: 120 W

**Parameters power** – The power [W]

Returns

This CalculatedTophatHeatSource object

# set\_scanning\_speed(beam\_speed: float = 0.5)

Sets the moving velocity of the heat source.

Default: 500.0e-3 m/s

Parameters beam\_speed – The beam speed [m/s]

#### Returns

This CalculatedTophatHeatSource object

# set\_wave\_length(wave\_length: float = 1064)

#### Parameters

wave\_length - The wavelength to be set for the heat source, defaults to 1064 nm.

### Returns

This CalculatedTophatHeatSource objectT

### with\_keyhole\_model(config: KeyholeModel)

Sets the keyhole model applied in the simulation.

# **Default: None**

Parameters config – The keyhole model

#### Returns

This CalculatedTophatHeatSource object

#### class tc\_python.am.CalibratedConicalHeatSource(java\_calibrated\_heat\_source)

Bases: object

A calibrated conical heat source.

### get\_absorptivity() $\rightarrow$ float

Returns the absorptivity of the heat source.

### Returns

The absorptivity [%]

 $\texttt{get\_hi}() \to \texttt{float}$ 

Returns the Hi dimension of the heat source.

#### Returns

The Hi dimension [micrometer]

# $\texttt{get\_re()} \to \texttt{float}$

Returns the Re dimension of the heat source.

### Returns

The Re dimension [micrometer]

# $\texttt{get\_ri()} \to \texttt{float}$

Returns the Ri dimension of the heat source.

# Returns

The Ri dimension [micrometer]

# $\texttt{get\_type()} \rightarrow \textit{HeatSourceType}$

Returns the type of heat source.

Returns The type

### set\_power(power: float = 120.0)

Sets the power of the heat source.

### Default: 120 W

Parameters power – The power [W]

### Returns

This CalibratedConicalHeatSource object

### set\_scanning\_speed(beam\_speed: float = 0.5)

Sets the moving velocity of the heat source.

Default: 500.0e-3 m/s

### Parameters

**beam\_speed** – The beam speed [m/s]

#### Returns

This CalibratedConicalHeatSource object

### class tc\_python.am.CalibratedDoubleEllipsoidalHeatSource(java\_calibrated\_heat\_source)

Bases: object

A calibrated double-ellipsoidal heat source.

#### get\_absorptivity() → float

Returns the absorptivity of the heat source.

#### Returns

The absorptivity [%]

### $\texttt{get\_af()} \rightarrow \texttt{float}$

Returns the Af dimension of the heat source.

### Returns

The Af dimension [micrometer]

## $\texttt{get\_ar()} \to \texttt{float}$

Returns the Ar dimension of the heat source.

#### Returns

The Ar dimension [micrometer]

# $\texttt{get\_b()} \to \texttt{float}$

Returns the B dimension of the heat source.

# Returns

The B dimension [micrometer]

# $\texttt{get\_c()} \to \texttt{float}$

Returns the C dimension of the heat source.

Returns

The C dimension [micrometer]

# $\texttt{get\_type()} \rightarrow \textit{HeatSourceType}$

Returns the type of heat source.

Returns The type

### set\_power(power: float = 120.0)

Sets the power of the heat source.

### Default: 120 W

Parameters power – The power [W]

### Returns

This CalibratedConicalHeatSource object

### set\_scanning\_speed(beam\_speed: float = 0.5)

Sets the moving velocity of the heat source.

Default: 500.0e-3 m/s

#### Parameters

**beam\_speed** – The beam speed [m/s]

#### Returns

This CalibratedDoubleEllipsoidalHeatSource object

#### class tc\_python.am.CalibratedGaussianHeatSource(java\_calibrated\_heat\_source)

Bases: object

A calibrated Gaussian heat source.

#### get\_absorptivity() → float

Returns the absorptivity of the heat source.

#### Returns

The absorptivity [%]

# get\_absorptivity\_prefactor() $\rightarrow$ float

Returns the absorptivity prefactor value of the calibrated heat source.

Returns

The absorptivity [%]

 $\texttt{get\_beam\_radius()} \rightarrow \texttt{float}$ 

Returns the beam radius of the heat source.

#### Returns

The beam radius [micrometer]

**get\_type()** → *HeatSourceType* 

Returns the type of heat source.

### Returns

The type

### $set_power(power: float = 120.0)$

Sets the power of the heat source.

Default: 120 W

Parameters

 $\textbf{power}-The \ power \ [W]$ 

### Returns

This CalibratedGaussianHeatSource object

# set\_scanning\_speed(beam\_speed: float = 0.5)

Sets the moving velocity of the heat source.

Default: 500.0e-3 m/s

# Parameters beam\_speed – The beam speed [m/s]

#### Returns

This CalibratedGaussianHeatSource object

#### class tc\_python.am.CalibratedHeatSource

Bases: HeatSource

A calibrated heat source.

# abstract set\_power(power: float = 120.0)

Sets the power of the heat source.

Default: 120 W

Parameters power – The power [W]

#### Returns

This CalibratedHeatSource object

abstract set\_scanning\_speed(beam\_speed: float = 0.5)

Sets the moving velocity of the heat source.

Default: 500.0e-3 m/s

# Parameters beam\_speed – The beam speed [m/s]

#### Returns

This CalibratedHeatSource object

#### class tc\_python.am.CoarseMesh

Bases: Mesh

An initially coarse mesh.

Note: It is adaptive and will be automatically refined as required.

## class tc\_python.am.ConicalHeatSource

Bases: object

A conical heat source.

#### $get_type() \rightarrow str$

Returns the type of heat source.

Returns

The type

# set\_absorptivity(absorptivity: float = 60.0)

Sets the absorptivity.

Default: 60%

Parametersabsorptivity – The absorptivity [%]

Returns

This ConicalHeatSource object

### **set\_hi**(*hi\_dim: float* = 0.0001)

Sets the parameter Hi that defines the dimensions of the heat source.

Tip: See Thermo-Calc Online Help for details about this heat source model.

**Default**: 100.0e-6 m

Parameters hi\_dim – The Hi parameter [m]

### Returns

This ConicalHeatSource object

# set\_power(power: float = 120.0)

Sets the power of the heat source.

Default: 120 W

Parameters power – The power [W]

### Returns

This ConicalHeatSource object

# **set\_re**(*re\_dim: float* = 0.0001)

Sets the parameter Re that defines the dimensions of the heat source.

Tip: See Thermo-Calc Online Help for details about this heat source model.

Default: 100.0e-6 m

Parameters re\_dim – The Re parameter [m]

Returns This ConicalHeatSource object

### **set\_ri**(*ri\_dim: float* = 6*e*-05)

Sets the parameter Ri that defines the dimensions of the heat source.

Tip: See Thermo-Calc Online Help for details about this heat source model.

Default: 60.0e-6 m

Parameters ri\_dim – The Ri parameter [m]

Returns

This ConicalHeatSource object

set\_scanning\_speed(beam\_speed: float = 0.5)

Sets the moving velocity of the heat source.

Default: 500.0e-3 m/s

Parameters beam\_speed – The beam speed [m/s]

Returns

This ConicalHeatSource object

**class** tc\_python.am.**CoreRingHeatSource**(*absorptivity: float* = 60.0)

Bases: object

A CoreRing heat source.

# disable\_keyhole\_model()

Disables using a keyhole model in the simulation.

# Returns

This CoreRingHeatSource object

### $\texttt{get\_type()} \to str$

Returns the type of heat source.

### Returns

The type

# set\_absorptivity(absorptivity: float = 60.0)

Sets the absorptivity as a constant.

Default: 60%

**Parameters absorptivity** – The absorptivity [%]

Returns

This CoreRingHeatSource object

set\_beam\_radius\_core(beam\_radius\_core: float = 5e-05)

Sets the beam radius-Core.

Default: 50.0e-6 m

Parameters beam\_radius\_core – The beam radius of the core beam [m]

Returns

This CoreRingHeatSource object

set\_beam\_radius\_ring(beam\_radius\_ring: float = 3.3e-05)

Sets the beam radius-Ring.

Default: 33.0e-6 m

**Note:** set\_beam\_radius\_ring() cannot be used with set\_core\_ring\_index().

**Parameters beam\_radius\_ring** – The beam radius of the ring beam [m]

#### Returns

This CoreRingHeatSource object

### set\_core\_ring\_index(core\_ring\_index: CoreRingIndex = CoreRingIndex.INDEX\_0)

Sets the *Core-Ring Index. Beam radius-Ring, Ring radius* and *Percentage of power in ring* are then set to a value based on the selected Index and Beam radius-Core.

**Default**: CoreRingIndex.INDEX\_0

**Note:** set\_core\_ring\_index() cannot be used with set\_beam\_radius\_ring(),
set\_ring\_radius() or set\_ring\_power\_percent().

#### **Parameters**

core\_ring\_index - The index (INDEX\_0 - INDEX\_6) of the Core-Ring laser system

#### Returns

This CoreRingHeatSource object

#### set\_power(power: float = 120.0)

Sets the power of the heat source.

Default: 120 W

Parameters power – The power [W]

#### Returns

This CoreRingHeatSource object

# set\_ring\_power\_percent(ring\_power\_percent: float = 7.0)

Sets the percentage of the power of the heat source in the ring beam

**Default**: 7.0 %

Note: set\_ring\_power\_percent() cannot be used with set\_core\_ring\_index.

# Parameters

**ring\_power\_percent** – The percentage of power in the ring bean [%]

#### Returns

This CoreRingHeatSource object

### set\_ring\_radius(ring\_radius: float = 9.858e-05)

Sets the ring radius.

Default: 98.58e-6 m

**Note:** set\_ring\_radius() cannot be used with set\_core\_ring\_index().

#### **Parameters**

ring\_radius – The radius of the ring beam [m]

#### Returns

This CoreRingHeatSource object

# set\_scanning\_speed(beam\_speed: float = 0.5)

Sets the moving velocity of the heat source.

Default: 500.0e-3 m/s

### Parameters beam\_speed – The beam speed [m/s]

#### Returns

This CoreRingHeatSource object

# with\_keyhole\_model(config: KeyholeModel)

Sets the keyhole model applied in the simulation.

# Default: None

#### **Parameters**

**config** – The keyhole model

### Returns

This CoreRingHeatSource object

# class tc\_python.am.CoreRingIndex(value)

# Bases: Enum

```
Index number of the Core-ring heat source. The selected Index sets a pre-defined value of Percentage of power in ring. INDEX_0 = 7%, INDEX_1 = 25%, INDEX_2 = 35%, INDEX_3 = 50%, INDEX_4 = 70%, INDEX_5 = 80%, INDEX_6 = 90%
```

### $INDEX_0 = 0$

*Percentage of power in ring* = 7%

# $INDEX_1 = 1$

*Percentage of power in ring* = 25%

# $INDEX_2 = 2$

*Percentage of power in ring* = 35%

# **INDEX\_3** = 3

*Percentage of power in ring* = 50%

# $INDEX_4 = 4$

*Percentage of power in ring* = 70%

# $INDEX_5 = 5$

*Percentage of power in ring* = 80%

### $INDEX_6 = 6$

*Percentage of power in ring* = 90%

**class** tc\_python.am.**CustomMesh**(*minimum\_element\_size: float = 1e-05, maximum\_element\_size: float = 0.0001*)

# Bases: Mesh

An initial mesh with explicitly defined dimensions.

Note: It is adaptive and will be automatically refined as required.

#### **Parameters**

- **minimum\_element\_size** The minimum element size [m]
- maximum\_element\_size The maximum element size [m]

tc\_python.am.DEFAULT\_CAMERA = {'position': [-2, -2, 1], 'viewup': [0, 0, 1]}

The default pyvista camera view.

# class tc\_python.am.DoubleEllipsoidalHeatSource

Bases: object

A double ellipsoidal heat source.

#### $get_type() \rightarrow str$

Returns the type of heat source.

Returns The type

#### set\_absorptivity(absorptivity: float = 60.0)

Sets the absorptivity.

Default: 60%

Parameters absorptivity – The absorptivity [%]

#### Returns

This DoubleEllipsoidalHeatSource object

# $set_af(af: float = 7e-05)$

Sets the parameter Af that defines the dimensions of the heat source.

Tip: See Thermo-Calc Online Help for details about this heat source model.

Default: 70.0e-6 m

**Parameters af** – The Af parameter [m]

Returns

This DoubleEllipsoidalHeatSource object

### $set_ar(ar: float = 70.0)$

Sets the parameter Ar that defines the dimensions of the heat source.

Tip: See Thermo-Calc Online Help for details about this heat source model.

Default: 70.0e-6 m

**Parameters ar** – The Ar parameter [m]

Returns

This DoubleEllipsoidalHeatSource object

#### **set\_b**(*b: float* = 8.5*e*-05)

Sets the parameter B that defines the dimensions of the heat source.

Tip: See Thermo-Calc Online Help for details about this heat source model.

### Default: 85.0e-6 m

**Parameters b** – The B parameter [m]

#### Returns

This DoubleEllipsoidalHeatSource object

### **set\_c**(*c*: *float* = 0.0002)

Sets the parameter C that defines the dimensions of the heat source.

Tip: See Thermo-Calc Online Help for details about this heat source model.

# Default: 200.0e-6 m

Parameters c – The C parameter [m]

# Returns

This DoubleEllipsoidalHeatSource object

 $set_power(power: float = 120.0)$ 

Sets the power of the heat source.

Default: 120 W

Parameters power – The power [W]

### Returns

This DoubleEllipsoidalHeatSource object

# set\_scanning\_speed(beam\_speed: float = 0.5)

Sets the moving velocity of the heat source.

Default: 500.0e-3 m/s

Parameters beam\_speed – The beam speed [m/s]

### Returns

This DoubleEllipsoidalHeatSource object

# class tc\_python.am.EveryNthTimeStep

Bases: FileSavingStrategy

Saving at every n-th time step.

### $\texttt{get\_type()} \to str$

Returns the type of the file saving strategy.

#### Returns

The type

# **set\_n**(*n*: *int* = 1)

Sets at which n-th time step files are saved.

### Default: 1

Parameters n – The n-th time step where files are saved

### Returns

This FileSavingStrategy object

### class tc\_python.am.EveryTimeInterval

Bases: FileSavingStrategy

Saving after regular time intervals.

# $\texttt{get\_type()} \to str$

Returns the type of the file saving strategy.

# Returns

The type

# set\_time\_interval(time\_interval: float = 0.01)

Sets the time interval at which files are saved.

Default: 0.01 s

**Parameters** time\_interval – The time interval [s]

#### Returns

This FileSavingStrategy object

### class tc\_python.am.FileSavingStrategy

Bases: object

The strategy for how result files are saved on disk. Both the number and time point of saving can be controlled.

### classmethod automatic()

An automatic saving strategy. This is the default.

### classmethod every\_n\_th\_time\_step()

Saving at every n-th time step.

# classmethod every\_time\_interval()

Saving after regular time intervals.

#### class tc\_python.am.FineMesh

Bases: Mesh

An initially fine mesh.

Note: It is adaptive and will be automatically refined as required.

# **class** tc\_python.am.**GaussianHeatSource**(*absorptivity: float* = 60.0)

Bases: object

A Gaussian heat source.

# disable\_keyhole\_model()

Disables using a keyhole model in the simulation.

### Returns

This GaussianHeatSource object

### $\texttt{get\_type()} \to \mathsf{str}$

Returns the type of heat source.

### Returns

The type

# set\_absorptivity(absorptivity: float = 60.0)

Sets the absorptivity as a constant.

Default: 60%

### Parameters absorptivity – The absorptivity [%]

#### Returns

This GaussianHeatSource object

# set\_beam\_radius(beam\_radius: float = 0.00011)

Sets the beam radius.

Default: 110.0e-6 m

#### Parameters

beam\_radius – The beam radius [m]

#### Returns

This GaussianHeatSource object

set\_power(power: float = 120.0)

Sets the power of the heat source.

Default: 120 W

# Parameters

 $\textbf{power}-The \ power \ [W]$ 

#### Returns

This GaussianHeatSource object

# set\_scanning\_speed(beam\_speed: float = 0.5)

Sets the moving velocity of the heat source.

### Default: 500.0e-3 m/s

Parameters beam\_speed – The beam speed [m/s]

### Returns

This GaussianHeatSource object

# with\_keyhole\_model(config: KeyholeModel)

Sets the keyhole model applied in the simulation.

# **Default: None**

Parameters config – The keyhole model

### Returns

This GaussianHeatSource object

# class tc\_python.am.HeatSource

Bases: object

The heat source.

The heat source model has either a Gaussian, Core-ring, Tophat, double ellipsoidal or conical distribution.

### Default: A Gaussian heat source.

# classmethod conical()

A conical heat source.

# The default is a Gaussian heat source.

# Returns

A new ConicalHeatSource object

**classmethod conical\_from\_library**(*heat\_source\_name: str, power: float = 120.0, beam\_speed: float = 0.5*)

Read a conical heat source given its name.

**Tip:** The available heat sources can be obtained using *HeatSource*. *get\_available\_calibrated\_heat\_source\_names()*. Their types are then available with the method get\_type() for each heat source.

#### **Parameters**

- heat\_source\_name The name of the conical heat source to read
- power The beam power, default: 120 W [W]
- beam\_speed The beam speed, default: 500e-3 m/s [m/s]

#### Returns

A new CalibratedConicalHeatSource object

```
classmethod core_ring_with_calculated_absorptivity(absorptivity_factor: float = 1.0,
```

wave\_length: float = 1064.0)

A core\_ring heat source with calculated absorptivity.

#### Parameters

- **absorptivity\_factor** A factor that adjusts the absorptivity of the heat source. Default is 1.0.
- wave\_length The wavelength is used for calculating absorptivity. Default is 1064.0 nm.

#### Returns

A new CalculatedCoreRingHeatSource object

#### classmethod core\_ring\_with\_constant\_absorptivity(absorptivity: float = 60.0)

A core\_ring heat source with constant absorptivity.

**Parameters absorptivity** – A constant value for absorptivity. Default is 60.0 %.

#### Returns

A new CoreRngHeatSource object

classmethod core\_ring\_with\_user\_defined\_function\_absorptivity(function: str = '60')

A core\_ring heat source with user defined function for absorptivity.

#### Parameters

**function** – The user defined function string for absorptivity, which can be constant, or a function of temperature (T), e.g. -1.6e-7\*T\*T+2.5e-3\*T+31.

#### Returns

A new UserDefinedCoreRingHeatSource object

#### classmethod double\_ellipsoidal()

A double ellipsoidal heat source.

# The default is a Gaussian heat source.

#### Returns

A new DoubleEllipsoidalHeatSource object

Read a double-ellipsoidal heat source given its name.

**Tip:** The available heat sources can be obtained using *HeatSource*. *get\_available\_calibrated\_heat\_source\_names()*. Their types are then available with the method get\_type() for each heat source.

### **Parameters**

- heat\_source\_name The name of the double-ellipsoidal heat source to read
- power The beam power, default: 120 W [W]
- beam\_speed The beam speed, default: 500e-3 m/s [m/s]

#### Returns

A new CalibratedDoubleEllipsoidalHeatSource object

**classmethod from\_library**(*heat\_source\_name: str, power: float = 120.0, beam\_speed: float = 0.5*) Read a heat source given its name.

**Tip:** The available heat sources can be obtained using *HeatSource*. *get\_available\_calibrated\_heat\_source\_names()*. Their types are then available with the method get\_type() for each heat source.

### **Parameters**

- heat\_source\_name The name of the heat source to read
- power The beam power, default: 120 W [W]
- beam\_speed The beam speed, default: 500e-3 m/s [m/s]

#### Returns

A new CalibratedHeatSource object

#### classmethod gaussian()

A Gaussian heat source.

#### Returns

A new GaussianHeatSource object

```
classmethod gaussian_from_library(heat_source_name: str, power: float = 120.0, beam_speed: float
= 0.5)
```

Read a Gaussian heat source given its name.

**Tip:** The available heat sources can be obtained using *HeatSource. get\_available\_calibrated\_heat\_source\_names()*. Their types are then available with the method get\_type() for each heat source.

# Parameters

- heat\_source\_name The name of the Gaussian heat source to read
- power The beam power, default: 120 W [W]
- beam\_speed The beam speed, default: 500e-3 m/s [m/s]

#### Returns

A new CalibratedGaussianHeatSource object

classmethod gaussian\_with\_calculated\_absorptivity(absorptivity\_factor: float = 1.0, wave\_length: float = 1064.0)

A gaussian heat source with calculated absorptivity.

#### Parameters

- **absorptivity\_factor** A factor that adjusts the absorptivity of the heat source. Default is 1.0.
- wave\_length The wavelength is used for calculating absorptivity. Default is 1064.0 nm.

#### Returns

A new CalculatedGaussianHeatSource object

# classmethod gaussian\_with\_constant\_absorptivity(absorptivity: float = 60.0)

A gaussian heat source with constant absorptivity.

#### Parameters

**absorptivity** – A constant value for absorptivity. Default is 60.0 %.

#### Returns

A new GaussianHeatSource object

# classmethod gaussian\_with\_user\_defined\_function\_absorptivity(function: str = '60')

A gaussian heat source with user defined function for absorptivity.

#### **Parameters**

**function** – The user defined function string for absorptivity, which can be constant, or a function of temperature (T), e.g. -1.6e-7\*T\*T+2.5e-3\*T+31.

#### Returns

A new UserDefinedGaussianHeatSource object

#### classmethod get\_available\_calibrated\_heat\_source\_names() $\rightarrow$ List[str]

Retrieves a list of available heat source names.

Tip: The heat sources can then be read using HeatSource.from\_library().

#### Returns

A list of strings representing the names of available heat sources

### $\texttt{classmethod get\_calibrated\_heatsource\_names()} \rightarrow List[str]$

Returns a list of the names of the available calibrated heat sources.

The name can then be used with the function get\_path\_of\_calibrated\_heatsource().

#### Returns

The list of names

### classmethod get\_path\_of\_calibrated\_heatsource( $optimized_heatsource_name: str$ ) $\rightarrow$ str

Returns the file path of the calibrated heat source.

### Parameters

optimized\_heatsource\_name – The name of the calibrated heat source

#### Returns

The file path

A tophat heat source with calculated absorptivity.

#### **Parameters**

- **absorptivity\_factor** A factor that adjusts the absorptivity of the heat source. Default is 1.0.
- wave\_length The wavelength is used for calculating absorptivity. Default is 1064.0 nm.

#### Returns

A new CalculatedCoreRingHeatSource object

# classmethod tophat\_with\_constant\_absorptivity(absorptivity: float = 60.0)

A tophat heat source with constant absorptivity.

#### Parameters

**absorptivity** – A constant value for absorptivity. Default is 60.0 %.

#### Returns

A new TophatHeatSource object

# classmethod tophat\_with\_user\_defined\_function\_absorptivity(function: str = '60')

A tophat heat source with user defined function for absorptivity.

#### **Parameters**

**function** – The user defined function string for absorptivity, which can be constant, or a function of temperature (T), e.g. -1.6e-7\*T\*T+2.5e-3\*T+31.

#### Returns

A new UserDefinedTophatHeatSource object

#### class tc\_python.am.HeatSourceType(value)

Bases: Enum

An enumeration.

CONICAL = 2

### DOUBLE\_ELLIPSOIDAL = 1

### GAUSSIAN = 0

### class tc\_python.am.KeyholeModel

Bases: object

A model for an "analytic" keyhole in the AM calculation.

# set\_beam\_quality(beam\_quality: float = 1.0)

Sets the Beam quality factor M2.

**Default**: 1.0

Parameters beam\_quality – The Beam quality factor M2 []

#### Returns

This KeyholeModel object

# set\_rayleigh\_length(rayleigh\_length: float = 0.0025)

Sets the Rayleigh length.

Default: 2.5e-3 m

# Parameters rayleigh\_length – The Rayleigh length [m]

#### Returns

This KeyholeModel object

# class tc\_python.am.LibraryMaterialProperties(library\_name: str)

Bases: MaterialProperties

Material properties previously saved on disk using the specified library name.

### class tc\_python.am.MaterialProperties

Bases: object

The material properties used in the AM calculation, can be either from a Scheil calculation or from a previously stored library.

### delete\_library()

Deletes the material library from disk.

# classmethod from\_library(library\_name: str)

Uses material properties previously saved on disk using a library name.

Parameters library\_name – The library name

#### Returns

A new LibraryMaterialProperties object

classmethod from\_scheil\_result(result: ScheilCalculationResult, interface\_scattering\_constant: float = 4e-08)

Creates material properties from the result of a Scheil calculation.

#### **Parameters**

- result The Scheil result to create material properties from
- interface\_scattering\_constant Greater than zero to use the interface scattering

# Returns

A new ScheilMaterialProperties object

### classmethod get\_all\_library\_names() $\rightarrow$ List[str]

Returns a list with the names of all material libraries available on disk.

### Returns

A list with the names of all material libraries

Zone, *to\_zone*: Zone)  $\rightarrow$  float

Returns average values for the specified material property in the specified zone interval.

#### **Parameters**

- material\_property\_enum The material property
- **from\_zone** The lower zone boundary

to\_zone – The upper zone boundary

#### Returns

The average value for the specified material property in the specified zone interval

#### get\_evaporation\_temperature() $\rightarrow$ float

Returns the evaporation temperature for the material.

#### Returns

The evaporation temperature [K]

#### get\_liquidus\_temperature() $\rightarrow$ float

Returns the liquidus temperature for the material.

#### Returns

The liquidus temperature [K]

# $get_name() \rightarrow str$

Returns the name of the library.

#### Returns

The name of the library

# $\texttt{get\_smoothed\_values\_for\_material\_property}(\textit{material\_property\_enum}: MaterialProperty}) \rightarrow \texttt{MaterialProperty}(material\_property\_enum) \rightarrow \texttt{MaterialProperty}(materialProperty) \rightarrow \texttt{MaterialProperty}(materialProperty\_enum) \rightarrow \texttt{MaterialProperty}(materialProperty\_enum) \rightarrow \texttt{MaterialProperty}(materialProperty\_enum) \rightarrow \texttt{MaterialProperty}(materialProperty) \rightarrow \texttt{MaterialProperty}(materialProperty\_enum) \rightarrow \texttt{MaterialProperty$

[List[float], List[float]]

Returns smoothed values for the specified material property.

#### Parameters

material\_property\_enum - The material property

#### Returns

The temperature [K] and the values of the specified material property

#### get\_smoothing\_for(material\_property\_enum: MaterialProperty) → Smoothing

Returns the smoothing level for the specified material property.

#### Parameters

material\_property\_enum - The material property to get smoothing level for

#### Returns

The smoothing level for the specified material property

### get\_solidification\_temperature() $\rightarrow$ float

Returns the solidification temperature for the material.

#### Returns

The solidification temperature [K]

### rename\_as\_library(name: str)

Renames the material library.

# Parameters

name – The new name of the library

### Returns

This MaterialProperties object

#### save\_as\_library(name: str = ")

Saves the material library with the specified name to disk.

Default: Re-save the current object with the previously chosen name

Parameters name – The new name of the library

#### Returns

This MaterialProperties object

# save\_library()

Saves the material library to disk.

#### Returns

This MaterialProperties object

### set\_smoothing\_for\_all\_properties(smoothing\_enum: Smoothing)

Sets the smoothing level for all material properties.

Parameters smoothing\_enum – The smoothing level

# Returns

This MaterialProperties object

set\_smoothing\_for\_property(material\_property\_enum: MaterialProperty, smoothing\_enum: Smoothing)

Sets the smoothing level for the specified material property.

### Parameters

- material\_property\_enum The specified material property
- **smoothing\_enum** The smoothing level

#### Returns

This MaterialProperties object

# class tc\_python.am.MaterialProperty(value)

# Bases: Enum

A single material property used in the class MaterialProperties.

# CP = 0

Apparent heat capacity [J/(kg K)]

### DENSITY = 1

Density [kg/m3]

### DRIVING\_FORCE\_EVAPORATION = 8

Driving force for evaporation [J/mol]

# DYNAMIC\_VISCOSITY = 4

Dynamic viscosity [Pa s]

# ELECTRIC\_RESISTIVITY = 11

Resistivity [Ohm \* m]

# ENTHALPY\_PER\_KG = 10 Enthalpy [J/kg]

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# ENTHALPY\_PER\_MOLE = 2

Enthalpy [J/mol]

# EVAPORATION\_ENTHALPY = 9

Evaporation enthalpy [J/mol]

# $MOLAR_MASS_OF_GAS = 7$

Molar mass of Gas [kg/mol]

# $MOLAR_VOLUME = 6$

Molar volume [m3/mol]

# SURFACE\_TENSION = 5

Surface tension [J/m2]

# THERMAL\_CONDUCTIVITY = 3

Thermal conductivity [W/(m K)]

# class tc\_python.am.MaterialType(value)

Bases: Enum

The material (solid, liquid, powder) to be plotted in a *pyvista* visualization plot.

## LIQUID = 'Liquid'

Onl liquid material

POWDER = 'Powder'

Only powder

# SHOW\_ALL = 'All'

All material

# SOLID = 'Solid'

Only solid material

### SOLID\_AND\_LIQUID = 'Solid and liquid'

Only solid and liquid material

# SOLID\_AND\_POWDER = 'Solid and powder'

Only solid material and powder

# class tc\_python.am.MediumMesh

Bases: Mesh

An initially medium mesh.

Note: It is adaptive and will be automatically refined as required.

# class tc\_python.am.Mesh

Bases: object

The initial mesh size in the simulation.

Can be coarse, medium, fine, or custom.

Note: It is adaptive and will be automatically refined as required.

### classmethod coarse()

Selecting the mesh to be initially coarse.

Note: It is adaptive and will be automatically refined as required.

### Returns

A new CoarseMesh object

**classmethod custom**(*minimum\_element\_size: float = 1e-05, maximum\_element\_size: float = 0.0001*) Selecting explicitly the initial mesh.

Default: Minimum element size: 10 um, maximum element size: 100 um

**Note:** It is adaptive and will be automatically refined as required.

#### **Parameters**

- minimum\_element\_size The minimum element size [m]
- maximum\_element\_size The maximum element size [m]

#### Returns

A new CustomMesh object

#### classmethod fine()

Selecting the mesh to be initially fine.

Note: It is adaptive and will be automatically refined as required.

#### Returns

A new FineMesh object

# classmethod medium()

Selecting the mesh to be initially medium.

Note: It is adaptive and will be automatically refined as required.

# Returns

A new MediumMesh object

### class tc\_python.am.NumericalOptions

Bases: object

The numerical options for an AM simulation.

# disable\_damping()

Disable numerical damping.

# **Default: disabled**
#### Returns

This NumericalOptions object

#### disable\_petrov\_galerkin()

Disables Streamline upwind Petrov-Galerkin (SUPG) for the numerical solver.

#### Default: enabled

#### Returns

This NumericalOptions object

#### enable\_petrov\_galerkin()

Enables Streamline upwind Petrov-Galerkin (SUPG) for the numerical solver.

#### **Default: enabled**

#### Returns

This NumericalOptions object

# get\_options()

Returns the numerical options for an AM simulation.

#### Returns

The numerical options

# set\_damping\_factor(damping\_factor: float = 0.0)

Sets the numerical damping factor.

# Default: numerical damping is disabled

#### **Parameters**

damping\_factor – The numerical damping factor

#### Returns

This NumericalOptions object

#### set\_number\_of\_cores(num\_cores: int)

Sets the number of used processor cores.

# Default: Half of the available cores on the CPU, 2 cores on a 2-core CPU, and 1 core on a 1-core CPU.

# Parameters

num\_cores – The number of used cores

#### Returns

This NumericalOptions object

#### set\_smagorinsky\_constant(const: float = 0.18)

Sets the smagorinsky constant.

#### Parameters

const – The smagorinsky constant.

#### Returns

This NumericalOptions object

#### with\_file\_saving\_strategy(file\_saving\_strategy: FileSavingStrategy)

Sets the strategy how result files are saved on disk. Both the number and time point of saving can be controlled.

#### Default: an automatic file saving strategy

Parameters file\_saving\_strategy – The file saving strategy

Returns

This NumericalOptions object

Bases: object

The coordinates of a probe. This is a point in the simulation domain whose properties can be obtained from the result object after the calculation using *TransientResult.get\_temperatures\_at\_probe()*.

#### class tc\_python.am.Scalar(value)

Bases: Enum

A quantity to be plotted in a *pyvista* visualization plot.

#### MATERIAL\_TYPE = 'subdomain\_id'

Material type

#### MOLAR\_VOLUME = 'molar\_volume'

Molar volume

```
SURFACE_TENSION = 'gamma'
```

Surface tension

# TEMPERATURE = 'temperature'

Temperature

# THERMAL\_CONDUCTIVITY = 'k'

Thermal conductivity

# VOLUME\_FRACTION\_LIQUID = 'liquid\_vfrac'

Volume fraction of liquid

# class tc\_python.am.ScanningPath(calc: AdditiveManufacturingCalculation)

Bases: object

# $\texttt{get\_end\_of\_scanning\_time()} \to \texttt{float}$

# Returns

The end of scanning time (s)

#### get\_heat\_source\_position(*time: float*) → Tuple[List[float], float]

#### **Parameters**

 $\texttt{time} - The \ time \ (s)$ 

#### Returns

position of the heat source, and its angle. Returns an empty position list, if the heat source is OFF

get\_heat\_source\_position\_nearby(*layerID: int, x: float, y: float, y: float*)  $\rightarrow$  Tuple[List[float], float, float]

#### **Parameters**

- **layerID** The layer ID. Zero index
- $\mathbf{x}$  x-coordinate of the point.
- **y** y-coordinate of the point.

#### Returns

position of the nearest point, factor of this position on the line, distance to the line, and time.

# **Parameters**

- **pathID** The path ID of the scanning path. Zero index
- layerID The layer ID of the scanning path. Zero index
- **factor** The factor. 0 for the beginning position, and 1- for the end position of the line path.

#### Returns

position of the heat source, its angle, and time.

```
\texttt{get\_number\_of\_layers()} \rightarrow \text{int}
```

#### Returns

Total number of layers

get\_position\_between\_two\_nearest\_lines(layerID: int, x: float, y: float) → Tuple[float]

# **Parameters**

- **layerID** The layer ID. Zero index
- $\mathbf{x}$  x-coordinate of the point.
- **y** y-coordinate of the point.

#### Returns

The nearest position in the middle of two scanning paths. The scanning paths are determined by the two closet to [x, y]

# get\_scanning\_path\_of\_layer(layer: int) → List[List[float]]

# Returns

List of scanning paths on a layer

# **Parameters**

- pathID The path ID of the scanning path. Zero index
- **layerID** The layer ID of the scanning path. Zero index

#### Returns

Stating time, and ending time of a path, on a layer.

#### $is\_heat\_source\_on(time: float) \rightarrow bool$

**Parameters** time – The time (s)

#### Returns

True if the heat source is ON

#### class tc\_python.am.ScanningStrategy

Bases: object

The scanning pattern of the heat source.

Single track, bidirectional (flipping scanning direction of the heat source between alternate tracks), or unidirectional (same scanning direction of the heat source for all tracks) are available.

# classmethod bi\_directional()

A bidirectional scanning strategy (flipping scanning direction of the heat source between alternate tracks).

#### Returns

A new BiDirectionalScanningStrategy object

# set\_cooling\_time(cooling\_time: float)

Sets the cooling time.

Default: 0 [s]

Parameters cooling\_time - [s]

#### Returns

This ScanningStrategy object

### classmethod single\_track()

A single track scanning strategy.

#### Returns

A new SingleTrackScanningStrategy object

# classmethod uni\_directional()

A unidirectional scanning strategy (same scanning direction of the heat source for all tracks).

# Returns

A new UniDirectionalScanningStrategy object

#### class tc\_python.am.ScheilMaterialProperties(scheil\_result: ScheilCalculationResult,

*interface\_scattering\_constant: float = 0.0, note: str = ")* 

Bases: MaterialProperties

Material properties created from the result of a Scheil calculation.

# class tc\_python.am.SingleTrackScanningStrategy

Bases: ScanningStrategy

A single track scanning strategy.

# $\texttt{get\_type()} \to \mathsf{str}$

Returns the type of scanning strategy.

# Returns

The type

### set\_margin(margin: float = 0.001)

Sets the margin.

This is the offset of the scanning path from the sides of the computational domain. In case of single tracks, offset is placed from the sides transverse to the scanning direction.

Default: 1.0e-3 [m]

Parameters margin – The margin [m]

Returns

This SingleTrackScanningStrategy object

set\_number\_of\_layers(number\_of\_layers: int = 1)

Sets the number of layers.

Default: 1

Parameters number\_of\_layers – The number of layers

Returns

This SingleTrackScanningStrategy object

set\_powder\_fill\_time(powder\_fill\_time: float = 0.0)

Sets the powder fill time.

Default: 0 [s]

Parameters powder\_fill\_time - [s]

Returns

This SingleTrackScanningStrategy object

class tc\_python.am.Smoothing(value)

Bases: Enum

The smoothing level used in the class MaterialProperties.

CONSTANT = -200

Constant smoothing

# LARGE = 600

Large smoothing

LINEAR = -100

Linear smoothing

LITTLE = 60

Little smoothing

# MEDIUM = 150

Medium smoothing

### NONE = 0

No smoothing

class tc\_python.am.SteadyStateCalculation(calculation)

Bases: AdditiveManufacturingCalculation

A steady-state Additive Manufacturing calculation.

**Note:** This computes the temperature distribution in a steady-state environment, either on a bare metal substrate or with a powder layer on the top, with the possibility to add fluid flow inside the melt pool.

#### $calculate(timeout_in_minutes: float = 0.0) \rightarrow SteadyStateResult$

Runs the calculation.

# Parameters

**timeout\_in\_minutes** – Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a :class`UnrecoverableCalculationException` will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

#### Returns

A *SteadyStateResult* which later can be used to get specific values from the calculated result

# class tc\_python.am.SteadyStateResult(result)

Bases: AdditiveManufacturingResult

A result for a steady-state calculation.

# $\texttt{get\_heat\_affected\_zone\_depth()} \rightarrow \texttt{float}$

Returns the depth of the heat affected zone.

#### Returns

The depth of the heat affected zone [m]

# $\texttt{get\_heat\_affected\_zone\_length()} \rightarrow \texttt{float}$

Returns the length of the heat affected zone.

#### Returns

The heat affected zone length [m]

# $\texttt{get\_heat\_affected\_zone\_width()} \rightarrow \texttt{float}$

Returns the width of the heat affected zone.

# Returns

The width of the heat affected zone [m]

#### get\_keyhole\_depth() → float

The depth of the keyhole.

#### Returns

The depth of the keyhole [m]

# $\texttt{get\_keyhole\_length()} \rightarrow \texttt{float}$

The length of the keyhole.

#### Returns

The length of the keyhole [m]

# $\texttt{get\_keyhole\_width()} \rightarrow \texttt{float}$

The width of the keyhole.

#### Returns

The width of the keyhole [m]

#### $get_meltpool_depth() \rightarrow float$

Returns the meltpool depth.

# Returns

The meltpool depth [m]

 $\texttt{get\_meltpool\_length()} \to \texttt{float}$ 

Returns the meltpool length.

Returns

The meltpool length [m]

 $\texttt{get\_meltpool\_width()} \rightarrow \texttt{float}$ 

Returns the meltpool width.

# Returns

The meltpool width [m]

 $has_keyhole() \rightarrow bool$ 

Returns if the result contains a keyhole.

# Returns

True if the result has a keyhole

# class tc\_python.am.TopBoundaryConditions

Bases: object

The top boundary conditions of the simulation.

# disable\_evaporation()

Disables the evaporation heat loss due to heating of the powder layer or the metallic surface when being close to the evaporation temperature.

# Default: enabled

### Returns

This TopBoundaryConditions object

# enable\_evaporation()

Enables the evaporation heat loss due to heating of the powder layer or the metallic surface when being close to the evaporation temperature.

Default: enabled

# Returns

This TopBoundaryConditions object

### set\_convective\_heat\_coefficient(convective\_heat\_coefficient: float = 20.0)

Sets the convective heat transfer coefficient for the top surface to the surrounding gas.

Enter 0 to disable convective heat transfer.

Default: 20.0 W/m\*\*2

### Parameters

convective\_heat\_coefficient – The convective heat transfer coefficient [W/m\*\*2]

### Returns

This TopBoundaryConditions object

# set\_radiation\_emissivity(radiation\_emissivity: float = 0.8)

Sets the radiation from the top surface to the surrounding gas.

Enter 0 to disable radiation.

Default: 0.8

# Parameters

radiation\_emissivity – The radiation emissivity, range: [0 - 1] [-]

Returns This TopBoundaryConditions object **class** tc\_python.am.**TophatHeatSource**(*absorptivity: float* = 60.0) Bases: object A Tophat heat source. disable\_keyhole\_model() Disables using a keyhole model in the simulation. Returns This TophatHeatSource object  $get_type() \rightarrow str$ Returns the type of heat source. Returns The type set\_absorptivity(absorptivity: float = 60.0) Sets the absorptivity as a constant. Default: 60% **Parameters absorptivity** – The absorptivity [%] Returns This TophatHeatSource object set\_beam\_radius(beam\_radius: float = 0.00011) Sets the beam radius. Default: 110.0e-6 m **Parameters beam\_radius** – The beam radius [m] Returns This TophatHeatSource object set\_power(power: float = 120.0) Sets the power of the heat source. Default: 120 W **Parameters power** – The power [W] Returns This TophatHeatSource object set\_scanning\_speed(beam\_speed: float = 0.5) Sets the moving velocity of the heat source. Default: 500.0e-3 m/s

Parameters

**beam\_speed** – The beam speed [m/s]

### Returns

This TophatHeatSource object

#### with\_keyhole\_model(config: KeyholeModel)

Sets the keyhole model applied in the simulation.

**Default:** None

Parameters config – The keyhole model

#### Returns

This TophatHeatSource object

# class tc\_python.am.TransientCalculation(calculation)

Bases: AdditiveManufacturingCalculation

A transient Additive Manufacturing calculation.

**Note:** This computes the temperature distribution in a transient case with the given scanning strategy including multiple paths and layers and the possibility to add fluid flow inside the melt pool.

#### add\_probe(coordinate: ProbeCoordinate)

Adds a probe, this a point in the simulation domain whose properties can be obtained from the result after the calculation using *TransientResult.get\_temperatures\_at\_probe()*.

#### **Parameters**

coordinate – The probe to be added

#### Returns

This TransientCalculation object

#### $calculate(timeout_in_minutes: float = 0.0) \rightarrow TransientResult$

Runs the calculation.

#### Parameters

**timeout\_in\_minutes** – Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

# Returns

A TransientResult which later can be used to get specific values from the calculated result

# get\_scanning\_path()

#### Returns

The calculation's scanning pattern object

# remove\_all\_probes()

Removes all probes.

# Returns

This TransientCalculation object

#### remove\_probe(coordinate: ProbeCoordinate)

Removes a probe.

#### Parameters

coordinate – The coordinates of the probe to be removed

#### Returns

This TransientCalculation object

#### set\_length(length: float = 0.005)

Sets the length of the simulation domain.

Default: 5.0e-3 m :param length: The length [m] :return: This TransientCalculation object

#### set\_width(width: float = 0.004)

Sets the width of the simulation domain.

#### Default: 4.0e-3 m

**Parameters** width – The width [m]

#### Returns

This TransientCalculation object

#### with\_scanning\_strategy(scanning\_strategy: ScanningStrategy)

Sets the scanning strategy of the heat source, i.e. beam.

# Parameters scanning\_strategy - The scanning strategy

Returns

This TransientCalculation object

# class tc\_python.am.TransientResult(result)

Bases: AdditiveManufacturingResult

A result for transient calculations (also with steady-state heat source).

# $get\_temperatures\_at\_probe(coordinate: ProbeCoordinate) \rightarrow Tuple[List[float], List[float]]$

Obtains the temperature at a probe (i.e., a point in the simulation domain) that had previously been defined for the calculation using *TransientCalculation.add\_probe()* or *TransientWithSteadyStateCalculation.add\_probe()*.

#### Parameters

**coordinate** – The coordinates of the probe - must have been defined in the calculator previously

#### Returns

A tuple (time points [s], temperatures [K])

### $\texttt{get\_time\_steps()} \rightarrow \texttt{List[float]}$

#### Returns

List of the available time steps

set\_active\_time(step: int)

Set the step to be active

# class tc\_python.am.TransientWithSteadyStateCalculation(transient\_with\_ss\_calculation,

ss calculation)

# Bases: AdditiveManufacturingCalculation

A transient Additive Manufacturing calculation using a heat source from steady-state.

**Note:** This computes the temperature distribution in a transient case with the given scanning strategy including multiple paths and layers.

1. A steady-state simulation runs with the configured heat source and with the possibility to add fluid flow in the melt pool.

2. A volume heat source (based on the solution of steady-state calculation) is used in the transient simulation.

**Tip:** This type of calculation is significantly faster than fully transient calculations using *TransientCalculation*.

#### add\_probe(coordinate: ProbeCoordinate)

Adds a probe, this a point in the simulation domain whose properties can be obtained from the result after the calculation using *TransientResult.get\_temperatures\_at\_probe()*.

Parameters coordinate – The probe to be added

#### Returns

This TransientWithSteadyStateCalculation object

 $calculate(timeout_in_minutes: float = 0.0) \rightarrow TransientResult$ 

Runs the calculation.

#### Parameters

**timeout\_in\_minutes** – Used to prevent the calculation from running longer than what is wanted, or from hanging. If the calculation runs longer than timeout\_in\_minutes, a UnrecoverableCalculationException will be thrown, the current TCPython-block will be unusable and a new TCPython block must be created for further calculations.

#### Returns

A TransientResult which later can be used to get specific values from the calculated result

# disable\_fluid\_flow\_marangoni()

Disables the fluid flow modelling of the Marangoni effect.

Default: Enabled

Returns

This TransientWithSteadyStateCalculation object

# disable\_separate\_materials()

Disables separate material properties for powder and solid material.

Default: Disabled

#### Returns

This TransientWithSteadyStateCalculation object

#### enable\_fluid\_flow\_marangoni()

Enables the fluid flow modelling of the Marangoni effect.

Default: Enabled

**Note:** This option is not possible to use in conjunction with the option *separate material*, which is therefore automatically disabled.

#### Returns

This TransientWithSteadyStateCalculation object

# enable\_separate\_materials()

Enables separate material properties for powder and solid material.

Default: Disabled

**Note:** This option is not possible to use in conjunction with the option *Marangoni fluid flow*, which is therefore automatically disabled.

#### Returns

This TransientWithSteadyStateCalculation object

# remove\_all\_probes()

Removes all probes.

#### Returns

This TransientWithSteadyStateCalculation object

#### remove\_probe(coordinate: ProbeCoordinate)

Removes a probe.

# Parameters

coordinate – The coordinates of the probe to be removed

#### Returns

This TransientWithSteadyStateCalculation object

# set\_ambient\_temperature(temperature: float = 296.15)

Sets the ambient temperature.

Default: 23 degree Celsius

# Parameters

temperature – The ambient temperature [K]

#### Returns

This TransientWithSteadyStateCalculation object

#### set\_base\_plate\_temperature(temperature: float = 303.15)

Sets the baseplate temperature.

Default: 30 degree Celsius

# Parameters

temperature – The baseplate temperature [K]

#### Returns

This TransientWithSteadyStateCalculation object

# set\_gas\_pressure(pressure: float = 100000.0)

Sets the gas pressure.

Default: 1.0e5 Pa

# Parameters pressure – The pressure [Pa]

#### Returns

This TransientWithSteadyStateCalculation object

set\_height(height: float = 0.002)

Sets the height of the simulation domain.

Default: 2.0e-3 m

**Parameters height** – The height [m]

# Returns

This TransientWithSteadyStateCalculation object

#### set\_layer\_thickness(layer\_thickness: float = 4e-05)

Sets the layer thickness.

Default: 40.0e-6 m

Parameters layer\_thickness – The layer thickness [m]

#### Returns

This TransientWithSteadyStateCalculation object

#### set\_length(length: float = 0.005)

Sets the length of the simulation domain.

**Default**: 5.0e-3 m :param length: The length [m] :return: This *TransientWithSteadyStateCalculation* object

# set\_powder\_density(powder\_density: float = 80.0)

Sets the powder density.

**Default:** 80.0%

Parameters powder\_density – The powder density [%]

### Returns

This TransientWithSteadyStateCalculation object

# set\_width(width: float = 0.004)

Sets the width of the simulation domain.

Default: 4.0e-3 m

Parameters width – The width [m]

#### Returns

This TransientWithSteadyStateCalculation object

# with\_heat\_source(heat\_source: HeatSource)

Sets the heat source.

# Parameters

heat\_source – The heat source

#### Returns

This TransientWithSteadyStateCalculation object

#### with\_material\_properties(material\_properties: MaterialProperties)

Sets the material properties.

**Tip:** Material properties can be defined like this: MaterialProperties.from\_library("IN718") or MaterialProperties.from\_scheil\_result(scheil\_result).

Parameters	
material_propert	<b>ies</b> – The material properties

#### Returns

This TransientWithSteadyStateCalculation object

with\_mesh(mesh: Mesh)

Sets the mesh.

Parameters mesh – The mesh

#### Returns

This TransientWithSteadyStateCalculation object

# with\_numerical\_options(numerical\_options: NumericalOptions)

Sets the numerical options.

Parameters numerical\_options – The numerical options

# Returns

This TransientWithSteadyStateCalculation object

# with\_scanning\_strategy(scanning\_strategy: ScanningStrategy)

Sets the scanning strategy of the heat source, i.e. beam.

# Parameters scanning\_strategy - The scanning strategy

#### Returns

This TransientWithSteadyStateCalculation object

#### with\_top\_boundary\_conditions(boundary\_conditions: TopBoundaryConditions)

Sets the boundary conditions.

# Parameters

boundary\_conditions – The boundary conditions

#### Returns

This TransientWithSteadyStateCalculation object

# class tc\_python.am.UniDirectionalScanningStrategy

Bases: ScanningStrategy

A unidirectional scanning strategy (same scanning direction of the heat source for all tracks).

# $\texttt{get\_type()} \to \mathsf{str}$

Returns the type of scanning strategy.

# Returns

The type

# set\_angle(angle: float = 0.0)

Sets the rotation of the scanning direction between two consecutive layers.

Note: The scanning direction of the first layer is always aligned parallel to the x-axis.

Default: 0 degree

Parameters angle – The angle [degree]

Returns

This UniDirectionalScanningStrategy object

# set\_hatch\_spacing(hatch\_spacing: float = 0.0)

Sets the horizontal separation between two consecutive tracks.

Default: 0 m

**Parameters** hatch\_spacing – The hatch spacing [m]

Returns

This UniDirectionalScanningStrategy object

# set\_lift\_time(lift\_time: float = 0.0)

Sets the lift time, i.e. the time between two tracks where the heat source is inactive.

Default: 0 s

Parameters lift\_time – The lift time [s]

#### Returns

This UniDirectionalScanningStrategy object

# set\_margin(margin: float = 0.001)

Sets the margin.

This is the offset of the scanning path from the sides of the computational domain.

**Default**: 1.0e-3 [m]

Parameters

margin – The margin [m]

#### Returns

This UniDirectionalScanningStrategy object

set\_number\_of\_layers(number\_of\_layers: int = 1)

Sets the number of layers.

# Default: 1

Parameters number\_of\_layers – The number of layers

# Returns

This UniDirectionalScanningStrategy object

set\_powder\_fill\_time(powder\_fill\_time: float = 0.0)

Sets the powder fill time.

Default: 0 s

Parameters
 powder\_fill\_time - The powder fill time [s]

Returns

This UniDirectionalScanningStrategy object

# **class** tc\_python.am.**UserDefinedCoreRingHeatSource**(*function: str* = '60.0')

Bases: object

A CoreRing heat source (with user defined absorptivity)

#### disable\_keyhole\_model()

Disables using a keyhole model in the simulation.

#### Returns

This UserDefinedCoreRingHeatSource object

# static get\_type() $\rightarrow$ str

Returns the type of heat source.

Returns

The type

# set\_absorptivity\_function(function: str = '60')

# **Parameters**

**function** – The absorptivity user defined function value to set for the heat source, provided as a string.

# Returns

This UserDefinedCoreRingHeatSource object

set\_beam\_radius\_core(beam\_radius\_core: float = 5e-05)

Sets the beam radius-Core.

Default: 50.0e-6 m

Parameters

beam\_radius\_core – The beam radius of the core beam [m]

#### Returns

This CoreRingHeatSource object

# set\_beam\_radius\_ring(beam\_radius\_ring: float = 3.3e-05)

Sets the beam radius-Ring.

Default: 33.0e-6 m

**Note:** set\_beam\_radius\_ring() cannot be used with set\_core\_ring\_index().

#### **Parameters**

beam\_radius\_ring – The beam radius of the ring beam [m]

#### Returns

This CoreRingHeatSource object

set\_core\_ring\_index(core\_ring\_index: CoreRingIndex = CoreRingIndex.INDEX\_0)

Sets the *Core-Ring Index. Beam radius-Ring, Ring radius* and *Percentage of power in ring* are then set to a value based on the selected Index and Beam radius-Core.

**Default**: CoreRingIndex.INDEX\_0

**Note:** set\_core\_ring\_index() cannot be used with set\_beam\_radius\_ring(),
set\_ring\_radius() or set\_ring\_power\_percent().

#### Parameters

```
core_ring_index - The index (INDEX_0 - INDEX_6) of the Core-Ring laser system
```

#### Returns

This CoreRingHeatSource object

#### set\_power(power: float = 120.0)

Sets the power of the heat source.

# Default: 120 W

**Parameters power** – The power [W]

#### Returns

This UserDefinedCoreRingHeatSource object

#### set\_ring\_power\_percent(ring\_power\_percent: float = 7.0)

Sets the percentage of the power of the heat source in the ring beam

**Default**: 7.0 %

**Note:** set\_ring\_power\_percent() cannot be used with set\_core\_ring\_index().

Parameters

**ring\_power\_percent** – The percentage of power in the ring bean [%]

#### Returns

This CoreRingHeatSource object

#### set\_ring\_radius(ring\_radius: float = 9.858e-05)

Sets the ring radius.

Default: 98.58e-6 m

**Note:** set\_ring\_radius() cannot be used with set\_core\_ring\_index().

# Parameters

ring\_radius – The radius of the ring beam [m]

# Returns

This CoreRingHeatSource object

# set\_scanning\_speed(beam\_speed: float = 0.5)

Sets the moving velocity of the heat source.

Default: 500.0e-3 m/s

Parameters beam\_speed – The beam speed [m/s]

#### Returns

This UserDefinedCoreRingHeatSource object

# with\_keyhole\_model(config: KeyholeModel)

Sets the keyhole model applied in the simulation.

#### **Default: None**

Parameters config – The keyhole model

#### Returns

This UserDefinedCoreRingHeatSource object

#### **class** tc\_python.am.**UserDefinedGaussianHeatSource**(*function: str* = '60.0')

Bases: object

A Gaussian heat source (with user defined absorptivity)

#### disable\_keyhole\_model()

Disables using a keyhole model in the simulation.

#### Returns

This GaussianHeatSource object

# static get\_type() $\rightarrow$ str

Returns the type of heat source.

Returns

The type

# set\_absorptivity\_function(function: str = '60')

### Parameters

**function** – The absorptivity user defined function value to set for the heat source, provided as a string.

# Returns

This GaussianHeatSource object

# set\_beam\_radius(beam\_radius: float = 0.00011)

Sets the beam radius.

Default: 110.0e-6 m

# Parameters

beam\_radius – The beam radius [m]

# Returns

This GaussianHeatSource object

### set\_power(power: float = 120.0)

Sets the power of the heat source.

Default: 120 W

Parameters power – The power [W]

#### Returns

This GaussianHeatSource object

# set\_scanning\_speed(beam\_speed: float = 0.5)

Sets the moving velocity of the heat source.

Default: 500.0e-3 m/s

Parameters beam\_speed – The beam speed [m/s]

#### Returns

This GaussianHeatSource object

# with\_keyhole\_model(config: KeyholeModel)

Sets the keyhole model applied in the simulation.

# **Default: None**

Parameters config – The keyhole model

Returns This GaussianHeatSource object

# **class** tc\_python.am.**UserDefinedTophatHeatSource**(*function: str* = '60.0')

Bases: object

A Tophat heat source (with user defined absorptivity)

# disable\_keyhole\_model()

Disables using a keyhole model in the simulation.

#### Returns

This UserDefinedTophatHeatSource object

# $\texttt{static get\_type()} \rightarrow str$

Returns the type of heat source.

#### Returns

The type

#### set\_absorptivity\_function(function: str = '60')

# Parameters

**function** – The absorptivity user defined function value to set for the heat source, provided as a string.

# Returns

This UserDefinedTophatHeatSource object

#### set\_beam\_radius(beam\_radius: float = 0.00011)

Sets the beam radius.

# Default: 110.0e-6 m

Parameters beam\_radius – The beam radius [m]

#### Returns

This UserDefinedTophatHeatSource object

#### set\_power(power: float = 120.0)

Sets the power of the heat source.

Default: 120 W

# Parameters

power - The power [W]

#### Returns

This UserDefinedTophatHeatSource object

# set\_scanning\_speed(beam\_speed: float = 0.5)

Sets the moving velocity of the heat source.

### Default: 500.0e-3 m/s

Parameters beam\_speed – The beam speed [m/s]

# Returns

This UserDefinedTophatHeatSource object

# with\_keyhole\_model(config: KeyholeModel)

Sets the keyhole model applied in the simulation.

# Default: None

Parameters config – The keyhole model

#### Returns

This UserDefinedTophatHeatSource object

# class tc\_python.am.Zone(value)

# Bases: Enum

Zones to be used in combination with keyhole model and material properties.

# LIQUID = 2

Liquid zone

# MUSHY = 1

Mushy zone

# SOLID = 0

Solid zone

# VAPOR = 3

Vapor zone

# 6.2 Module "system"

# class tc\_python.system.MultiDatabaseSystemBuilder(multi\_database\_system\_builder)

Bases: object

Used to select databases, elements, phases etc. and create a System object. The difference to the class System-Builder is that the operations are performed on all the previously selected databases. The system is then used to create calculations.

# create\_and\_select\_species(stoichiometry: str)

Specify a species from the already entered elements. The stoichiometry of the species is the chemical formula of the species. The created species will also be automatically selected.

**Note:** The elements in the chemical formula are normally separated by stoichiometric numbers. Neither parenthesis "()" nor an underscore "\_" is allowed in the chemical formula, while the special combination "/-" or "/+" can be used. Consult the Thermo-Calc database documentation for details about the syntax.

# Parameters stoichiometry – The stoichiometry of the species

#### Returns

This MultiDatabaseSystemBuilder object

#### deselect\_constituent\_on\_sublattice(phase\_name: str, sublattice\_no: int,

constituent\_name\_to\_deselect: str)

Rejects a constituent on a sublattice in a phase in both the thermodynamic and the kinetic database.

#### **Parameters**

- phase\_name The name of the phase
- **sublattice\_no** The number of the sublattice (starting with 1)
- constituent\_name\_to\_deselect The name of the constituent to deselect

#### Returns

This MultiDatabaseSystemBuilder object

#### deselect\_phase(phase\_name\_to\_deselect: str)

Rejects a phase for both the thermodynamic and the kinetic database.

### **Parameters**

phase\_name\_to\_deselect - The phase name

#### Returns

This MultiDatabaseSystemBuilder object

#### deselect\_species(species\_name: str)

Removes the species from the system.

#### Parameters

**species\_name** – The species

#### Returns

This MultiDatabaseSystemBuilder object

#### $get_system() \rightarrow System$

Creates a new System object that is the basis for all calculation types. Several calculation types can be defined later from the object; these are independent.

#### Returns

A new System object

Selects a constituent on a sublattice in a phase in both the thermodynamic and the kinetic database.

**Note:** Previously the third parameter *constituent\_name\_to\_select* had a wrong name, it has been corrected in version 2021b.

#### **Parameters**

- **phase\_name** The name of the phase
- **sublattice\_no** The number of the sublattice (starting with 1)
- constituent\_name\_to\_select The name of the constituent to select

#### Returns

This MultiDatabaseSystemBuilder object

#### select\_phase(phase\_name\_to\_select: str)

Selects a phase for both the thermodynamic and the kinetic database.

# Parameters phase\_name\_to\_select – The phase name

Returns

This MultiDatabaseSystemBuilder object

#### select\_species(species\_name: str)

Adds the species to the system. Up to 1000 species can be defined in a single system.

# Parameters

species\_name - The species

#### Returns

This MultiDatabaseSystemBuilder object

#### with\_new\_composition\_set(composition\_set: CompositionSet)

Used to enter two or more composition sets for a phase. If a phase has a miscibility gap it is necessary to have two composition sets, one for each possible composition that can be stable simultaneously.

The databases often create the typical composition sets for phases automatically when data are retrieved. The equilibrium calculations (using the default settings with global minimization) will usually add new composition sets if needed.

**Note:** Precipitation and diffusion calculations can require the user to define additional composition sets. E.g. in the case where the new composition set is needed in the configuration of the calculation.

#### **Parameters**

composition\_set - the composition set

#### Returns

This MultiDatabaseSystemBuilder object

# without\_default\_phases()

Rejects all the default phases from both the thermodynamic and the kinetic database, any phase now needs to be selected manually for the databases.

#### Returns

This MultiDatabaseSystemBuilder object

class tc\_python.system.System(system\_instance)

Bases: object

A system containing selections for databases, elements, phases etc.

**Note:** For the defined system, different calculations can be configured and run. **Instances of this class should always be created from a SystemBuilder**.

**Note:** The system object is **immutable**, i.e. it cannot be changed after is has been created. If you want to change the system, you must instead create a new one.

**convert\_composition**(*input\_composition*: *Dict[str, float]*, *input\_unit*: ConversionUnit, *output\_unit*: ConversionUnit, *dependent\_component*: str = ")  $\rightarrow$  Dict[str, float]

Provides conversion between composition units for any combination of chemical compounds. It is fast because no thermodynamic equilibrium calculation is involved.

Syntax of the chemical compounds: "Al2O3", "FeO", "CO", "Fe", "C", ...

**Note:** It is not required that the chemical compounds are components of the database. The only requirement is that all elements are present in the database.

#### **Parameters**

- input\_composition Composition (for example: { "Al2O3": 25.0, "FeO": 75.0})
- input\_unit Unit of the input composition
- **output\_unit** Requested output unit
- dependent\_component The dependent component (optional), for example: "Fe". If no dependent component is specified the sum of the input composition needs to match 100% / 1

#### Returns

The composition in the requested output unit

```
get_all_elements_in_databases() \rightarrow List[str]
```

Returns the names of all elements present in the selected databases, regardless of the actual selection of elements.

#### Returns

A list of element names

#### $\texttt{get\_all\_phases\_in\_databases()} \rightarrow \texttt{List[str]}$

Returns all phase names present in the selected databases, regardless of selected elements, phases etc.

#### Returns

A list of phase names

# get\_all\_species\_in\_databases() $\rightarrow$ List[str]

Returns all species names present in the selected databases, regardless of the actual selection of elements, phases, ....

Returns

A list of species names

#### get\_database\_names() $\rightarrow$ List[str]

Returns the names of the selected thermodynamic and mobility databases.

#### Returns

A list of database names

#### get\_element\_object(element\_name: str) → Element

Returns the Element object of an element. This can be used to obtain detailed information about the element.

# Parameters element\_name – The element name

#### Returns

A Element: object

#### get\_elements\_in\_system() → List[str]

Returns the names of all elements present in the selected system.

**Note:** The list does not contain any elements or components that have been auto-selected by the database(s) in a calculator. Use the get\_components() of the calculator object instead to get the complete information.

#### Returns

A list of element names

# get\_phase\_object(phase\_name: str) → Phase

Returns the Phase object of a phase. This can be used to obtain detailed information about the phase.

# Parameters

phase\_name – The phase name

#### Returns

A Phase: object

# $\texttt{get\_phases\_in\_system()} \rightarrow List[str]$

Returns all phase names present in the system due to its configuration (selected elements, phases, etc.).

#### Returns

A list of phase names

# $\texttt{get\_references()} \rightarrow Dict[str, List[str]]$

Provides a dictionary with database references per database in the selected system.

#### Returns

The database references

# $\texttt{get\_species\_in\_system()} \rightarrow List[str]$

Returns the names of all species present in the selected system.

**Note:** The list does not contain any species or components that have been auto-selected by the database(s) in a calculator. Use the get\_components() of the calculator object instead to get the complete information.

#### Returns

The list of species names

### 

Returns the Species object of an species. This can be used to obtain detailed information about the species.

#### Parameters

species\_name – The species name

#### Returns

A Species: object

# $get_system_data() \rightarrow SystemData$

Returns the content of the database. This can be used to modify the parameters and functions and to change the current system by using with\_system\_modifications().

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as \*.tdb-file.

#### Returns

The system data

# with\_batch\_equilibrium\_calculation(default\_conditions: bool = True, components: List[str] = []) $\rightarrow$ BatchEquilibriumCalculation

Creates a batch-equilibrium calculation (a vectorized equilibrium calculation).

**Note:** Use this instead of looping if you want to calculate equilibria for a larger number of compositions and know the conditions in advance. This calculation type has improved performance when calculating a large number of equilibria when each individual calculations is quick. E.g. when evaluating single phase properties for thousands of compositions.

### Parameters

- default\_conditions If *True*, automatically sets the conditions *N*=1 and *P*=100000
- **components** Specify here the components of the system (for example: [AL2O3, ...]), *only necessary if they differ from the elements*. If this option is used, **all elements** of the system need to be replaced by a component.

#### Returns

A new BatchEquilibriumCalculation object

# with\_cct\_precipitation\_calculation() $\rightarrow$ PrecipitationCCTCalculation

Creates a CCT diagram calculation.

#### Returns

A new PrecipitationCCTCalculation object

#### with\_isothermal\_diffusion\_calculation() $\rightarrow$ DiffusionIsoThermalCalculation

Creates an isothermal diffusion calculation.

#### Returns

A new DiffusionIsoThermalCalculation object

#### with\_isothermal\_precipitation\_calculation() $\rightarrow$ PrecipitationIsoThermalCalculation

Creates an isothermal precipitation calculation.

#### Returns

A new PrecipitationIsoThermalCalculation object

with\_material\_to\_material()  $\rightarrow$  MaterialToMaterialCalculationContainer

Provides access to all Material to Material calculations. The actual calculation needs to be chosen in the returned object.

#### Returns

A new MaterialToMaterialCalculationContainer object

with\_non\_isothermal\_diffusion\_calculation()  $\rightarrow$  DiffusionNonIsoThermalCalculation

Creates a non-isothermal precipitation calculation.

#### Returns

A new PrecipitationNonIsoThermalCalculation object

with\_non\_isothermal\_precipitation\_calculation()  $\rightarrow$  *PrecipitationNonIsoThermalCalculation* 

Creates a non-isothermal precipitation calculation.

#### Returns

A new PrecipitationNonIsoThermalCalculation object

with\_phase\_diagram\_calculation(default\_conditions: bool = True, components:  $List[str] = []) \rightarrow PhaseDiagramCalculation$ 

Creates a phase diagram (map) calculation.

#### **Parameters**

- default\_conditions If *True*, automatically sets the conditions *N*=1 and *P*=100000
- **components** Specify here the components of the system (for example: [AL2O3, ...]), *only necessary if they differ from the elements*. If this option is used, **all elements** of the system need to be replaced by a component.

#### Returns

A new PhaseDiagramCalculation object

with\_property\_diagram\_calculation(default\_conditions: bool = True, components: List[str] = [])  $\rightarrow$ PropertyDiagramCalculation

Creates a property diagram (step) calculation.

#### Parameters

- default\_conditions If True, automatically sets the conditions N=1 and P=100000
- **components** Specify here the components of the system (for example: [AL2O3, ...]), *only necessary if they differ from the elements*. If this option is used, **all elements** of the system need to be replaced by a component.

#### Returns

A new PropertyDiagramCalculation object

```
with_property_model_calculation(model: str, path_to_models: str = ", debug_model: bool = False) \rightarrow
PropertyModelCalculation
```

Creates a Property Model calculation.

The parameter *debug\_model* is only used when debugging self-developed models.

### **Parameters**

- model The Property Model to be calculated.
- **path\_to\_models** The path where the Property Models are installed. If no value is entered, the Property Models folder used by the normal Thermo-Calc application is used.
- debug\_model Used when debugging self-developed models.

### Returns

A new PropertyModelCalculation object

# with\_scheil\_calculation() $\rightarrow$ ScheilCalculation

Creates a Scheil solidification calculation.

### Returns

A new ScheilCalculation object

with\_single\_equilibrium\_calculation(default\_conditions: bool = True, components: List[str] = [])  $\rightarrow$  SingleEquilibriumCalculation

Creates a single equilibrium calculation.

# Parameters

- default\_conditions If *True*, automatically sets the conditions *N*=1 and *P*=100000
- **components** Specify here the components of the system (for example: [AL2O3, ...]), *only necessary if they differ from the elements*. If this option is used, **all elements** of the system need to be replaced by a component.

### Returns

A new SingleEquilibriumCalculation object

# with\_ttt\_precipitation\_calculation() $\rightarrow$ PrecipitationTTTCalculation

Creates a TTT diagram calculation.

### Returns

A new PrecipitationTTTCalculation object

class tc\_python.system.SystemBuilder(system\_builder)

Bases: object

Used to select databases, elements, phases etc. and create a System object. The system is then used to create calculations.

# create\_and\_select\_species(stoichiometry: str)

Specify a species from the already entered elements. The stoichiometry of the species is the chemical formula of the species. The created species will also be automatically selected.

**Note:** The elements in the chemical formula are normally separated by stoichiometric numbers. Neither parenthesis "()" nor an underscore "\_" is allowed in the chemical formula, while the special combination "/-" or "/+" can be used. Consult the Thermo-Calc database documentation for details about the syntax.

Parameters stoichiometry – The stoichiometry of the species

#### Returns

This SystemBuilder object

Rejects a constituent on a sublattice in a phase in the last specified database only.

#### Parameters

- phase\_name The name of the phase
- **sublattice\_no** The number of the sublattice (starting with 1)
- constituent\_name\_to\_deselect The name of the constituent to deselect

#### Returns

This SystemBuilder object

# deselect\_phase(phase\_name\_to\_deselect: str)

Rejects a phase in the last specified database only.

#### **Parameters**

phase\_name\_to\_deselect - The name of the phase

#### Returns

This SystemBuilder object

#### deselect\_species(stoichiometry: str)

Removes the species from the system.

# Parameters stoichiometry – The species

#### Returns

This SystemBuilder object

# $get_system() \rightarrow System$

Creates a new System object that is the basis for all calculation types. Several calculation types can be defined later from the object; these are independent.

#### Returns

A new System object

Selects a constituent on a sublattice in a phase in the last specified database only.

**Note:** Previously the third parameter *constituent\_name\_to\_select* had a wrong name, it has been corrected in version 2021b.

#### **Parameters**

- **phase\_name** The name of the phase
- **sublattice\_no** The number of the sublattice (starting with 1)
- constituent\_name\_to\_select The name of the constituent to select

#### Returns

This SystemBuilder object

select\_database\_and\_elements(database\_name: str, list\_of\_element\_strings: List[str])

Selects a thermodynamic or kinetic database and its selected elements (that will be appended). After that, phases can be selected or unselected.

# Parameters

- database\_name The database name, for example "FEDEMO"
- list\_of\_element\_strings A list of one or more elements as strings, for example ["Fe", "C"]

# Returns

This SystemBuilder object

#### select\_phase(phase\_name\_to\_select: str)

Selects a phase in the last specified database only.

Parameters
 phase\_name\_to\_select - The name of the phase

#### Returns

This SystemBuilder object

#### select\_species(stoichiometry: str)

Adds the species to the system. Up to 1000 species can be defined in a single system.

# Parameters stoichiometry – The species

#### Returns

This SystemBuilder object

select\_user\_database\_and\_elements(path\_to\_user\_database: str, list\_of\_element\_strings: List[str])

Selects a thermodynamic database which is a user-defined database and select its elements (that will be appended).

**Note:** By using a r-literal, it is possible to use slashes on all platforms, also on Windows: *select\_user\_database\_and\_elements(r"my path/user\_db.tdb", ["Fe", "Cr"]])* 

Otherwise it is required to use **double** back-slashes on Windows as separator.

**Note:** On Linux and Mac the path is case-sensitive, also the file ending.

#### **Parameters**

- **path\_to\_user\_database** The path to the database file ("database".TDB), defaults to the current working directory. Only the filename is required if the database is located in the same folder as the script.
- list\_of\_element\_strings A list of one or more elements as strings, for example ["Fe", "C"]

#### Returns

This SystemBuilder object

#### with\_new\_composition\_set(composition\_set: CompositionSet)

Used to enter composition sets for a phase. If a phase has a miscibility gap it is necessary to have two composition sets, one for each possible composition that can be stable simultaneously.

#### Parameters

composition\_set - The composition set

#### Returns

This SystemBuilder object

# without\_default\_phases()

Rejects all default phases in the last specified database only, any phase needs now to be selected manually for that database.

Returns

This SystemBuilder object

# 6.3 Module "entities"

#### class tc\_python.entities.CompositionSet(phase\_name: str)

Bases: object

Used by the method tc\_python.system.SystemBuilder.with\_new\_composition\_set() to enter two or more composition sets for a phase.

#### Parameters

phase\_name - The name of the phase for which a new composition set is required

set\_major\_constituents\_for\_sublattice(sublattice\_index: int, major\_constituents: List[str])

Specify the new major constituent(s) for the sublattice.

Default: If not specified, a default is automatically chosen based on the specified composition set.

**Note:** This is useful in order to make calculations converge faster and more easily (because it may simplify giving start values when calculating the equilibrium as those phases with miscibility gaps should have different major constituents for each composition set). **The databases often set major constituents for several phases automatically when the data is retrieved.** 

#### **Parameters**

- **sublattice\_index** Index of the sublattice to set the major constituents for (starting with *I*)
- **major\_constituents** Optional list of the major constituents, which must be selected from the phase constitution of the current system.

#### Returns

This CompositionSet object

# class tc\_python.entities.Element(element)

# Bases: object

Represents an element, making detailed information about the element accessible.

### get\_enthalpy() $\rightarrow$ float

Returns the enthalpy of the element at 298 K, part of the stable element reference state (SER).

#### Returns

The enthalpy [J]

# get\_entropy\_diff\_0\_to\_298k() $\rightarrow$ float

Returns the entropy difference 0 - 298 K of the element, part of the stable element reference state (SER).

#### Returns

The entropy difference 0 - 298 K [J/K]

# $\texttt{get\_molar\_mass()} \to \texttt{float}$

Returns the molar mass of the element.

#### Returns

The molar mass [g/mol]

#### $get_name() \rightarrow str$

Returns the name of the element.

# Returns

The element name

# $\texttt{get\_stable\_element\_reference()} \rightarrow str$

Returns the stable element reference (i.e. the stable phase at 298.15 K and 1 bar, reference for all element thermodynamic data).

#### Returns

The name of the stable element reference

# $\texttt{is\_interstitial()} \rightarrow \texttt{bool}$

Returns if the element is interstitial.

**Note:** In the diffusion simulations (DICTRA), the assumption that the volume is carried by the substitutional elements only is applied. The interstitial elements are assumed to have zero molar volumes.

#### Returns

If the element is interstitial

# $\texttt{is\_special()} \rightarrow \texttt{bool}$

Returns if the element is special (i.e. vacancies (VA) and electrons (denoted either as /- in gaseous, liquid or solid phases, or ZE in an aqueous solution phase)).

# Returns

If the element is special

# $\texttt{is\_valid()} \rightarrow \texttt{bool}$

Returns if the element is valid. Non-valid elements are represented by an empty name.

#### Returns

If the element is valid

# class tc\_python.entities.Phase(phase)

Bases: object

Represents a phase, making detailed information about the phase accessible.

#### $get_name() \rightarrow str$

Returns the name of the phase.

# Returns

The phase name

 $get_species() \rightarrow Set[Species]$ 

Returns the species of the phase.

#### Returns

A set containing the species

# $\texttt{get\_species\_for\_composition\_profile()} \rightarrow \texttt{Set}[Species]$

Returns all species that need to be defined in a composition profile of the phase for diffusion simulations - except for one species that needs to be the dependent species.

**Note:** In a composition profile of a phase for diffusion simulations it is necessary to specify all non-stoichiometric and non-special species. In case of a DILUTE diffusion model, the database enforces the choice of a certain dependent species.

#### Returns

Set with the species

# get\_sublattices() $\rightarrow$ List[Sublattice]

Returns the sublattices of the phase in a well-defined contiguous order.

#### Returns

A list containing the Sublattice objects

# $get_type() \rightarrow PhaseType$

Returns the type of the phase (liquid, ionic liquid, solid, gas).

# Returns

The type of a phase

#### $has\_diffusion\_data() \rightarrow bool$

Returns if diffusion data exists for the phase.

# Returns

If diffusion data exists for the phase

# $\texttt{has\_molar\_volume\_data()} \rightarrow \texttt{bool}$

Returns if molar volume data exists for the phase.

# Returns

If molar volume data exists for the phase

# $\texttt{is_dilute_diffusion\_model()} \rightarrow \texttt{bool}$

Returns if diffusion is described using the DILUTE model for the phase. This will always return *False* if no diffusion data is available.

#### Returns

If the DILUTE model is used

# $\texttt{is\_gas()} \rightarrow \texttt{bool}$

Returns if the phase is a gas phase.

**Returns** If the phase is a gas phase

is\_ionic\_liquid()  $\rightarrow$  bool

Returns if the phase is an ionic liquid phase.

#### Returns

If the phase is an ionic liquid phase

# $\textbf{is\_liquid()} \rightarrow bool$

Returns if the phase is a liquid or ionic liquid phase.

# Returns

If the phase is a liquid phase

#### $\texttt{is\_solid}() \rightarrow \texttt{bool}$

Returns if the phase is a solid phase.

# Returns

If the phase is a solid phase

# class tc\_python.entities.PhaseType(value)

Bases: Enum

The type of a phase.

# GAS = 0

Gas phase.

#### IONIC\_LIQUID = 2

Ionic liquid phase.

# LIQUID = 1

Liquid phase.

# SOLID = 3

Solid phase.

# class tc\_python.entities.Species(species)

Bases: object

Represents a species, making detailed information about the species accessible.

# get\_all\_elements() → List[Tuple[*Element*, float]]

Returns all the elements that the species is composed of.

# Returns

List of all elements of the species and their stoichiometry

# $\texttt{get\_charge()} \rightarrow \texttt{int}$

Returns the charge of the species.

**Returns** The charge of the species

#### $get_name() \rightarrow str$

Returns the name of the species.

#### Returns

The species name

# $\texttt{is\_element()} \rightarrow \texttt{bool}$

Returns if the species actually represents an element.

# Returns

If the species represents an element

# $\texttt{is\_interstitial()} \rightarrow \texttt{bool}$

Returns if the species is interstitial.

**Note:** In the diffusion simulations (DICTRA), the assumption that the volume is carried by the substitutional elements only is applied. The interstitial elements are assumed to have zero molar volumes.

# Returns

If the species is interstitial

# $\texttt{is\_special()} \rightarrow \texttt{bool}$

Returns if the species is special (i.e. vacancies (VA) and electrons (denoted either as /- in gaseous, liquid or solid phases, or ZE in an aqueous solution phase)).

# Returns

If the species is special

# $\texttt{is\_valid()} \rightarrow \texttt{bool}$

Returns if the species is valid. Non-valid species are represented by an empty name.

# Returns

If the species is valid

# $\texttt{to\_element()} \rightarrow \textit{Element}$

Returns the *Element* representation of the species - if the species actually represents an element.

# Returns

The Element object

# class tc\_python.entities.Sublattice(sublattice)

Bases: object

Represents a sublattice of a phase.

# $get\_constituents() \rightarrow Set[Species]$

Returns the constituents of the sublattice.

### Returns

A set containing the constituents

# $\texttt{get\_nr\_of\_sites()} \to \texttt{float}$

Returns the number of sites in the sublattice.

# Returns

A float number

# 6.4 Module "server"

class tc\_python.server.AdditiveManufacturingCalculations(additive\_manufacturing\_calculations)
 Bases: object

Provides access to the calculation objects for all Additive Manufacturing calculations.

with\_steady\_state\_calculation(moose\_commands: Optional[Dict[str, str]] = None)  $\rightarrow$ SteadyStateCalculation

Creates a steady-state calculation for Additive Manufacturing.

**Note:** This computes the temperature distribution in a steady-state environment, either on a bare metal substrate or with a powder layer on the top, with the possibility to add fluid flow inside the melt pool.

**Warning:** It should not be necessary for most users to send commands directly to the FEM-solver, try to use the corresponding method implemented in the API instead.

#### Parameters

**moose\_commands** – Additional commands (key-value pairs) directly sent to the FEM solver - this is a special option that is normally not needed

Returns

A new SteadyStateCalculation object

with\_transient\_calculation(moose\_commands: Optional[Dict[str, str]] = None)  $\rightarrow$ TransientCalculation

Creates a transient calculation for Additive Manufacturing.

**Note:** This computes the temperature distribution in a transient case with the given scanning strategy including multiple paths and layers and the possibility to add fluid flow inside the melt pool.

**Warning:** It should not be necessary for most users to send commands directly to the FEM-solver, try to use the corresponding method implemented in the API instead.

# Parameters

**moose\_commands** – Additional commands (key-value pairs) directly sent to the FEM solver - this is a special option that is normally not needed

#### Returns

A new TransientCalculation object

with\_transient\_with\_steady\_state\_calculation(moose\_commands: Optional[Dict[str, str]] = None)  $\rightarrow$  TransientWithSteadyStateCalculation

Creates a transient calculation (with steady-state heat source) for Additive Manufacturing.

**Note:** This computes the temperature distribution in a transient case with the given scanning strategy including multiple paths and layers.

- 1. A steady-state simulation runs with the configured heat source and with the possibility to add fluid flow in the melt pool.
- 2. A volume heat source (based on the solution of steady-state calculation) is used in the transient simulation.

**Tip:** This type of calculation is significantly faster than fully transient calculations using TransientCalculation.

**Warning:** It should not be necessary for most users to send commands directly to the FEM-solver, try to use the corresponding method implemented in the API instead.

#### Parameters

**moose\_commands** – Additional commands (key-value pairs) directly sent to the FEM solver - this is a special option that is normally not needed

#### Returns

A new TransientWithSteadyStateCalculation object

# class tc\_python.server.LoggingPolicy(value)

Bases: Enum

Logging policy that determines how the TC-Python logs are presented to the user.

# FILE = 1

Logging to a file.

# NONE = 2

No logging at all.

# SCREEN = 0

Logging to the screen.

#### class tc\_python.server.MetallurgyCalculations(metallurgy\_calculations)

Bases: object

Provides access to the calculation objects for all Process Metallurgy calculations.

These are specialised calculations for working with metallurgical processes. Both equilibrium calculations and kinetic process simulations (Effective Equilibrium Reaction Zone model) are available.

with\_adiabatic\_equilibrium\_calculation(database: ProcessDatabase)  $\rightarrow$ AdiabaticEquilibriumCalculation

Creates an adiabatic equilibrium calculation for Process Metallurgy.

#### Parameters

database - The thermodynamic database used in the calculation

#### Returns

A new AdiabaticEquilibriumCalculation object

with\_adiabatic\_process\_calculation(database: ProcessDatabase)  $\rightarrow$  ProcessSimulationCalculation

Creates an adiabatic kinetic process simulation (EERZ, i.e. Effective Equilibrium Reaction Zone model).
Parameters

database – The thermodynamic database used in the calculation

Returns

A new ProcessSimulationCalculation object

with\_isothermal\_equilibrium\_calculation(database: ProcessDatabase) →

*IsoThermalEquilibriumCalculation* 

Creates an isothermal equilibrium calculation for Process Metallurgy.

#### Parameters

database – The thermodynamic database used in the calculation

Returns

A new IsoThermalEquilibriumCalculation object

#### class tc\_python.server.ResultLoader(result\_loader)

Bases: object

Contains methods for loading results from previously done calculations.

#### **diffusion**(*path: str*) $\rightarrow$ *DiffusionCalculationResult*

Loads a DiffusionCalculationResult from disc.

## **Parameters path** – path to the folder where result was previously saved.

#### Returns

A new DiffusionCalculationResult object which later can be used to get specific values from the calculated result

#### **phase\_diagram**(*path: str*) $\rightarrow$ *PhaseDiagramResult*

Loads a PhaseDiagramResult from disc.

#### Parameters

**path** – path to the folder where result was previously saved.

#### Returns

A new PhaseDiagramResult object which later can be used to get specific values from the calculated result

## $precipitation_TTT_or_CCT(path: str) \rightarrow PrecipitationCalculationTTTorCCTResult$

Loads a PrecipitationCalculationTTTorCCTResult from disc.

## **Parameters**

**path** – path to the folder where result was previously saved.

## Returns

A new PrecipitationCalculationTTTorCCTResult object which later can be used to get specific values from the calculated result

#### **precipitation\_single**(*path: str*) $\rightarrow$ *PrecipitationCalculationSingleResult*

Loads a PrecipitationCalculationSingleResult from disc.

#### Parameters

**path** – path to the folder where result was previously saved.

## Returns

A new PrecipitationCalculationSingleResult object which later can be used to get specific values from the calculated result

## **property\_diagram**(*path: str*) → *PropertyDiagramResult*

Loads a PropertyDiagramResult from disc.

## Parameters

**path** – path to the folder where result was previously saved.

#### Returns

A new PropertyDiagramResult object which later can be used to get specific values from the calculated result

## **property\_model**(*path: str*) → *PropertyModelResult*

Loads a PropertyModelResult from disc.

## Parameters

**path** – path to the folder where result was previously saved.

## Returns

A new PropertyModelResult object which later can be used to get specific values from the calculated result

## scheil(path: str) $\rightarrow$ ScheilCalculationResult

Loads a ScheilCalculationResult from disc.

#### Parameters

**path** – path to the folder where result was previously saved.

#### Returns

A new ScheilCalculationResult object which later can be used to get specific values from the calculated result

## 

Loads a SingleEquilibriumResult from disc.

## Parameters

**path** – path to the folder where result was previously saved.

#### Returns

A new SingleEquilibriumResult object which later can be used to get specific values from the calculated result

#### class tc\_python.server.SetUp(debug\_logging=False)

Bases: object

Starting point for all calculations.

Note: This class exposes methods that have no precondition, it is used for choosing databases and elements.

## disable\_caching()

A previously set cache folder is no longer used.

Note: Within the session, caching is activated and used through the default temporary directory.

#### Returns

This SetUp object

## 

Obtains the short information available for the specified database.

## Parameters

database\_short\_name – The name of the database (i.e. "FEDEMO", ...)

Returns

The short information about the database

## 

Obtains the path to the database file on disk. *TCPATH* is a placeholder for the root path of the used Thermo-Calc installation.

**Note:** Encrypted databases (\*.*TDC*) cannot be edited.

## **Parameters**

database\_short\_name – The name of the database (i.e. "FEDEMO", ...)

Returns

The path to the database on disk

## $\texttt{get\_databases()} \rightarrow List[str]$

Obtains the short names of all databases available in the used Thermo-Calc installation.

Note: Only databases with a valid license are listed.

## Returns

List of the available databases

## $get\_license\_manager() \rightarrow LicenseManager$

Returns the license manager, which is used for managing Thermo-calc licenses for TC-Python

#### Returns

the license manager

## 

Lists the names of all Property Models in the specified directory.

If the directory is not specified, the Property Model folder used by the normal Thermo-Calc application is used.

#### **Parameters**

**path\_to\_models** – The path where the Property Models are installed. If no value is entered, the Property Model folder used by the normal Thermo-Calc application is used.

#### Returns

Set containing all Property Model names

## $load_result_from_disk() \rightarrow ResultLoader$

Loads a previously calculated result from disk.

**Note:** This **only** works for results created by calling one of the **save\_result()** methods on a *Result* class created from a calculation.

## Returns

A new ResultLoader object

select\_database\_and\_elements(database\_name: str, list\_of\_elements: List[str])  $\rightarrow$  SystemBuilder Selects a first thermodynamic or kinetic database and selects the elements in it.

## Parameters

- database\_name The name of the database, for example "FEDEMO"
- **list\_of\_elements** The list of the selected elements in that database, for example ["Fe", "C"]

Returns

A new SystemBuilder object

*list\_of\_elements: List[str]*)  $\rightarrow$  *MultiDatabaseSystemBuilder* 

Selects the thermodynamic and kinetic database at once, guarantees that the databases are added in the correct order. Further rejection or selection of phases applies to both databases.

## **Parameters**

- thermodynamic\_db\_name The thermodynamic database name, for example "FEDEMO"
- kinetic\_db\_name The kinetic database name, for example "MFEDEMO"
- **list\_of\_elements** The list of the selected elements in that database, for example ["Fe", "C"]

## Returns

A new MultiDatabaseSystemBuilder object

# $select\_user\_database\_and\_elements(path\_to\_user\_database: str, list\_of\_elements: List[str]) \rightarrow SystemBuilder$

Selects a user-defined database and selects the elements in it.

**Note:** By using a r-literal, it is possible to use slashes on all platforms, also on Windows: *select\_user\_database\_and\_elements(r"my path/user\_db.tdb", ["Fe", "Cr"]])* 

Otherwise it is required to use **double** back-slashes on Windows as separator.

**Note:** On Linux and Mac the path is case-sensitive, also the file ending.

## **Parameters**

- **path\_to\_user\_database** The path to the database file ("database".TDB), defaults to the current working directory. Only filename is required if the database is located in the same folder as the script.
- **list\_of\_elements** The list of the selected elements in that database, for example ["Fe", "C"]

## Returns

A new SystemBuilder object

## set\_cache\_folder(path: str = ", precision\_for\_floats: int = 12)

Sets a folder where results from calculations and state of systems are saved. If at any time a calculation is run which has the exact same setting as a previous, the calculation is not re-run. The result is instead loaded from this folder.

**Note:** The same folder can be used in several scripts, and it can even be shared between different users. It can be a network folder.

## Parameters

- path path to the folder where results should be stored. It can be relative or absolute.
- **precision\_for\_floats** The number of significant figures used when comparing if the calculation has the same setting as a previous.

#### Returns

This SetUp object

### set\_ges\_version(version: int = 6)

Setting the version of the Gibbs Energy System (GES).

## Parameters version – The GES-version (currently version 5 or 6)

## Returns

This SetUp object

## set\_log\_level\_to\_debug()

Sets log level to DEBUG

## Returns

This SetUp object

## set\_log\_level\_to\_info()

Sets log level to INFO

#### Returns

This SetUp object

## with\_additive\_manufacturing() $\rightarrow$ AdditiveManufacturingCalculations

Provides access to the calculation objects for all Additive Manufacturing calculations.

## with\_metallurgy() $\rightarrow$ MetallurgyCalculations

Provides access to the calculation objects for all Process Metallurgy calculations.

These are specialised calculations for working with metallurgical processes. Both equilibrium calculations and kinetic process simulations (Effective Equilibrium Reaction Zone model) are available.

Bases: object

Starting point of the API. Typical syntax:

```
with TCPython() as session:
    session.select_database_and_elements(...)
```

**Note:** Each usage of *with TCPython()* causes significant overhead (starting a new process, stopping the old one, cleaning up the temporary disk space). Usually it is recommendable to call *with TCPython()* only once for each process, even if working in a loop. Instead you should pass the session or calculator object into the loop and use them there.

If necessary, beginning from version 2019a it is however possible to call with TCPython safely multiple times.

Starts a process of the API server and sets up the socket communication with it.

## Parameters

- **logging\_policy** Determines if the TC-Python log output is sent to the screen (*LoggingPolicy.SCREEN*), to file (*LoggingPolicy.FILE*) or nothing is logged at all (*LoggingPolicy.NONE*) **Default**: *LoggingPolicy.SCREEN*. Note that the log-handlers can also be adapted through the *tc\_python.LOGGER* object at any time.
- **log\_file** The log-file relative to the current path or absolute, only relevant if *log-ging\_policy=LoggingPolicy.FILE*. Log-output will be appended.
- **debug\_mode** If *True* it is tried to open a connection to an already running API-server. This is only used for debugging the API itself.
- is\_unittest Should be True if called by a unit test, only to be used internally for development.
- **do\_throw\_on\_backend\_hard\_crash** If *True* an *UnrecoverableCalculationException* will be thrown if the Java-backend crashes hard, if *False* the application will simply crash with a FORTRAN-stacktrace. If '*True*' the exception can be caught outside of the 'with'-clause and the application can continue, if '*False*' more information about the error is shown by the stacktrace.
- **port\_number** The port number for the communication with the Java-backend server. This is not required to be changed by normal users.

**Warning:** Most users should use *TCPython* using a with-statement for automatic management of the resources (network sockets and temporary files). If you anyway need to use that method, make sure to call *stop\_api\_server()* in any case using the try-finally-pattern.

tc\_python.server.stop\_api\_server(gateway\_id: Optional[str] = None)

Clears all resources used by the session (i.e. shuts down the API server and deletes all temporary files). The disk usage of temporary files might be significant.

**Warning:** Call this method only if you used *start\_api\_server()* initially. It should never be called when the API has been initialized in a with-statement using *TCPython*.

## 6.5 Module "quantity\_factory"

## class tc\_python.quantity\_factory.DiffusionQuantity

Bases: AbstractQuantity

Factory class providing quantities used for defining diffusion simulations and their results.

**Note:** In this factory class only the most common quantities are defined, you can always use the *Console Mode* syntax strings in the respective methods as an alternative (for example: "NPM(\*)").

**classmethod** activity\_of\_component(*component: str, use\_ser: bool* = *False*)  $\rightarrow$  ActivityOfComponent Creates a quantity representing the activity of a component.

## **Parameters**

- component The name of the component, use ALL\_COMPONENTS to choose all components
- **use\_ser** Use Stable-Element-Reference(SER). The user-defined reference state is be used if this setting is set to *False*.

## Returns

A new ActivityOfComponent object.

**classmethod chemical\_diffusion\_coefficient**(*phase: str, diffusing\_element: str, gradient\_element: str, reference\_element: str*)  $\rightarrow$ ChemicalDiffusionCoefficient

Creates a quantity representing the chemical diffusion coefficient of a phase [m^2/s].

## Parameters

- phase The name of the phase
- diffusing\_element The diffusing element
- gradient\_element The gradient element
- reference\_element The reference element (for example "Fe" in a steel)

## Returns

A new ChemicalDiffusionCoefficient object.

classmethod chemical\_potential\_of\_component(component: str, use\_ser: bool = False)  $\rightarrow$  ChemicalPotentialOfComponent

Creates a quantity representing the chemical potential of a component [J].

## Parameters

- **component** The name of the component, use *ALL\_COMPONENTS* to choose all components
- **use\_ser** Use Stable-Element-Reference(SER). The user-defined reference state is used if this setting is set to *False*.

## Returns

A new ChemicalPotentialOfComponent object.

**classmethod distance**(*region:* str = 'All')  $\rightarrow$  Distance

Creates a quantity representing the distance [m].

**Parameters region** – The name of the region or *All* to choose global.

classmethod intrinsic\_diffusion\_coefficient(phase: str, diffusing\_element: str, gradient\_element:

*str*, *reference\_element: str*)  $\rightarrow$  IntrinsicDiffusionCoefficient

Creates a quantity representing the intrinsic diffusion coefficient of a phase [m^2/s].

**Parameters** 

- **phase** The name of the phase
- diffusing\_element The diffusing element
- gradient\_element The gradient element
- reference\_element The reference element (for example "Fe" in a steel)

#### Returns

A new IntrinsicDiffusionCoefficient object.

**classmethod l\_bis**(*phase: str, diffusing\_element: str, gradient\_element: str, reference\_element: str*)  $\rightarrow$  Lbis

Creates a quantity representing L" of a phase [m^2/s].

## Parameters

- **phase** The name of the phase
- diffusing\_element The diffusing element
- gradient\_element The gradient element
- reference\_element The reference element (for example "Fe" in a steel)

#### Returns

A new Lbis object.

**classmethod mass\_fraction\_of\_a\_component**(*component: str*)  $\rightarrow$  MassFractionOfAComponent Creates a quantity representing the mass fraction of a component.

#### **Parameters**

**component** – The name of the component or ALL\_COMPONENTS to choose all components

Returns

A new MassFractionOfAComponent object.

**classmethod** mass\_fraction\_of\_a\_phase(*phase:* str)  $\rightarrow$  MassFractionOfAPhase

Creates a quantity representing the mass fraction of a phase.

## Parameters

phase – The name of the phase or ALL\_PHASES to choose all phases.

#### Returns

A new MassFractionOfAPhase object.

classmethod mobility\_of\_component\_in\_phase(phase: str, component: str) →

MobilityOfComponentInPhase

Creates a quantity representing the mobility of a component in a phase [m<sup>2</sup>/Js].

#### **Parameters**

- phase The name of the phase
- **component** The name of the component

#### Returns

A new MobilityOfComponentInPhase object.

classmethod mole\_fraction\_of\_a\_component(component: str)  $\rightarrow$  MoleFractionOfAComponent

Creates a quantity representing the mole fraction of a component.

## Parameters

component - The name of the component or ALL\_COMPONENTS to choose all components

#### Returns

A new MoleFractionOfAComponent object.

#### classmethod mole\_fraction\_of\_a\_phase(*phase: str*) $\rightarrow$ MoleFractionOfAPhase

Creates a quantity representing the mole fraction of a phase.

#### Parameters

phase – The name of the phase or ALL\_PHASES to choose all phases

#### Returns

A new MoleFractionOfAPhase object.

#### classmethod position\_of\_lower\_boundary\_of\_region(region: str) $\rightarrow$

PositionOfLowerBoundaryOfRegion

Creates a quantity representing the position of lower boundary of a region [m].

## Parameters

region – The name of the region

#### Returns

A new PositionOfLowerBoundaryOfRegion object.

## classmethod position\_of\_upper\_boundary\_of\_region(region: str) $\rightarrow$

PositionOfUpperBoundaryOfRegion

Creates a quantity representing the position of upper boundary of a region [m].

## Parameters

region – The name of the region

#### Returns

A new PositionOfUpperBoundaryOfRegion object.

## classmethod temperature() $\rightarrow$ Temperature

Creates a quantity representing the temperature [K].

#### Returns

A new Temperature object.

**classmethod thermodynamic\_factor**(*phase: str, diffusing\_element: str, gradient\_element: str, reference\_element: str*)  $\rightarrow$  ThermoDynamicFactor

Creates a quantity representing thermodynamic factor of a phase.

## Parameters

- **phase** The name of the phase
- diffusing\_element The diffusing element
- gradient\_element The gradient element
- reference\_element The reference element (for example "Fe" in a steel)

#### Returns

A new ThermoDynamicFactor object.

#### classmethod time() $\rightarrow$ Time

Creates a quantity representing the time [s].

### classmethod total\_mass\_fraction\_of\_component(component: str) →

TotalMassFractionOfComponent

Creates a quantity representing the total mass fraction of a component.

#### **Parameters**

**component** – The name of the component

Returns

A new TotalMassFractionOfComponent object.

#### classmethod total\_mass\_fraction\_of\_component\_in\_phase(phase: str, component: str) $\rightarrow$

TotalMassFractionOfComponentInPhase

Creates a quantity representing the total mass fraction of a component in a phase.

## **Parameters**

- **phase** The name of the phase
- component The name of the component

#### Returns

A new TotalMassFractionOfComponentInPhase object.

## **classmethod total\_mass\_fraction\_of\_phase**(*phase: str*) $\rightarrow$ TotalMassFractionOfPhase

Creates a quantity representing the total mass fraction of a phase.

## Parameters

**phase** – The name of the phase.

#### Returns

A new TotalMassFractionOfPhase object.

#### classmethod total\_mole\_fraction\_of\_component(component: str) →

TotalMoleFractionOfComponent

Creates a quantity representing the total mole fraction of a component.

#### Parameters

component – The name of the component

#### Returns

A new TotalMoleFractionOfComponent object.

 $\texttt{classmethod total_mole_fraction_of_component_in_phase}(\textit{phase: str, component: str}) \rightarrow \texttt{classmethod total_mole_fraction_of_component_in_phase}(\textit{phase: str, component: str_in_phase}(\textit{phase: str, component_in_phase}(\textit{phase: str, componen$ 

TotalMoleFractionOfComponentInPhase

Creates a quantity representing the total mole fraction of a component in a phase.

## Parameters

- **phase** The name of the phase
- component The name of the component

## Returns

A new TotalMoleFractionOfComponentInPhase object.

## **classmethod total\_volume\_fraction\_of\_phase**(*phase: str*) $\rightarrow$ TotalVolumeFractionOfPhase

Creates a quantity representing the total volume fraction of a phase.

#### **Parameters**

**phase** – The name of the phase.

## Returns

A new TotalVolumeFractionOfPhase object.

classmethod tracer\_diffusion\_coefficient(phase: str, diffusing\_element: str)  $\rightarrow$ TracerDiffusionCoefficient

Creates a quantity representing tracer diffusion coefficient of a phase [m^2/s].

## Parameters

• phase – The name of the phase

• diffusing\_element - The diffusing element

#### Returns

A new TracerDiffusionCoefficient object.

classmethod u\_fraction\_of\_a\_component(component: str)  $\rightarrow$  UFractionOfAComponent

Creates a quantity representing the u-fraction of a component.

Parameters

**component** – The name of the component

Returns

A new UFractionOfAComponent object.

## $\texttt{classmethod user\_defined\_function}(\textit{expression: str}) \rightarrow \texttt{Function}$

Creates a quantity representing a user-defined function.

## Parameters

**expression** – The function expression

#### Returns

A new Function object

## classmethod velocity\_of\_lower\_boundary\_of\_region(region: str) $\rightarrow$

VelocityOfLowerBoundaryOfRegion

Creates a quantity representing the velocity of lower boundary of a region [m/s].

#### Parameters

region – The name of the region

#### Returns

A new VelocityOfLowerBoundaryOfRegion object.

## classmethod velocity\_of\_upper\_boundary\_of\_region(region: str) $\rightarrow$

VelocityOfUpperBoundaryOfRegion

Creates a quantity representing the velocity of upper boundary of a region [m/s].

#### **Parameters**

region – The name of the region

#### Returns

A new VelocityOfUpperBoundaryOfRegion object.

## **classmethod width\_of\_region**(*region: str*) $\rightarrow$ Function

Creates a quantity representing the width of a region [m].

#### Parameters

region – The name of the region

#### Returns

A new WidthOfRegion object.

## class tc\_python.quantity\_factory.IndependentVariable

Bases: AbstractQuantity

Factory class providing quantities used for defining the independent variable in general diffusion result querying.

```
classmethod distance(region: str = 'All') \rightarrow Distance
```

Creates an independent variable representing the distance [m].

Returns

A new Distance object

```
<code>classmethod time()</code> \rightarrow <code>Time</code>
```

Creates an independent variable representing the time [s].

Returns

A new Time object

## class tc\_python.quantity\_factory.PlotCondition

Bases: AbstractQuantity

Factory class providing quantities used for defining the plot condition in general diffusion result querying.

**Note:** In this factory class only the most common quantities are defined, you can always use the *Console Mode* syntax strings in the respective methods as an alternative (for example: "time last").

```
classmethod distance(distancepoint: float, region: str = 'All') \rightarrow DistanceCondition
```

Creates a plot condition representing the distance [m].

Change in version 2019b: Mandatory parameter distancepoint added

## **Parameters**

- distancepoint The distance from the lower interface of the region
- region The name of the region or All to choose global.

#### Returns

A new DistanceCondition object

## $\texttt{classmethod integral()} \rightarrow IntegralCondition$

Creates an integral plot condition.

#### Returns

A new IntegralCondition object

**classmethod interface**(*region: str, interface\_position:* InterfacePosition)  $\rightarrow$  InterfaceCondition Creates a plot condition representing an interface between two regions.

#### **Parameters**

- region The name of the region used for defining the interface
- **interface\_position** The position of the interface relative to that region (lower or upper)

## Returns

A new InterfaceCondition object

**classmethod time**(*timepoint: Union*[*float, str*] = 'Last')  $\rightarrow$  TimeCondition

Creates a plot condition representing the time [s].

Change in version 2019b: Lists of timepoints are no longer supported

#### Parameters

timepoint - The timepoint. Optionally "Last" can be used for the end of the simulation

#### Returns

A new TimeCondition object

## class tc\_python.quantity\_factory.ScheilQuantity

## Bases: AbstractQuantity

Factory class providing quantities used for defining a Scheil calculation result (tc\_python.scheil. ScheilCalculationResult).

**classmethod** apparent\_heat\_capacity\_per\_gram()  $\rightarrow$  ApparentHeatCapacityPerGram

Creates a quantity representing the apparent heat capacity [J/g/K].

## Returns

A new ApparentHeatCapacityPerGram object.

 $\texttt{classmethod apparent\_heat\_capacity\_per\_mole()} \rightarrow ApparentHeatCapacityPerMole$ 

Creates a quantity representing the apparent heat capacity [J/mol/K].

#### Returns

A new ApparentHeatCapacityPerMole object.

## ${\tt classmethod\ apparent\_volumetric\_thermal\_expansion\_coefficient()} \rightarrow$

ApparentVolumetricThermalExpansionCoefficient

Creates a quantity representing the apparent volumetric thermal expansion coefficient of the system [1/K].

#### Returns

A new ApparentVolumetricThermalExpansionCoefficient object.

 $\texttt{classmethod average\_composition\_of\_solid\_phases\_as\_mass\_fraction(\textit{component: str}) \rightarrow \texttt{classmethod average\_composition\_component: str})$ 

AverageCompositionOfSolidPhasesAsMass

Creates a quantity representing the average composition of solid phases [mass fraction] at the current Scheil step.

## Parameters

**component** – The name of the component, use *ALL\_COMPONENTS* to choose all components

#### Returns

A new AverageCompositionOFSolidPhasesAsMassFraction object.

## $\texttt{classmethod average\_composition\_of\_solid\_phases\_as\_mole\_fraction(\textit{component: str}) \rightarrow \texttt{classmethod average\_composition\_component: str})$

AverageCompositionOfSolidPhasesAsMole

Creates a quantity representing the average composition of solid phases [mole fraction] at the current Scheil step.

## Parameters

**component** – The name of the component, use *ALL\_COMPONENTS* to choose all components

## Returns

A new AverageCompositionOFSolidPhasesAsMoleFraction object.

Creates a quantity representing the composition of a phase [mole-fraction].

## Parameters

- phase The name of the phase, use ALL\_PHASES to choose all stable phases
- component The name of the component, use ALL\_COMPONENTS to choose all components

#### Returns

A new CompositionOfPhaseAsMoleFraction object.

classmethod composition\_of\_phase\_as\_weight\_fraction(phase: str, component: str)  $\rightarrow$ 

CompositionOfPhaseAsWeightFraction

Creates a quantity representing the composition of a phase [weight-fraction].

#### **Parameters**

- phase The name of the phase, use ALL\_PHASES to choose all stable phases
- **component** The name of the component, use *ALL\_COMPONENTS* to choose all components

## Returns

A new CompositionOfPhaseAsWeightFraction object.

## classmethod density\_of\_phase(phase: str) $\rightarrow$ DensityOfPhase

Creates a quantity representing the average density of a phase [g/cm^3].

Parameters phase – The name of the phase or *ALL\_PHASES* to choose all phases

## Returns

A new DensityOfPhase object.

## $\texttt{classmethod density_of_system()} \rightarrow DensityOfSystem$

Creates a quantity representing the average density of the system [g/cm^3].

#### Returns

A new DensityOfSystem object.

 $\texttt{classmethod distribution_of_component_of_phase(\textit{phase: str, component: str})} \rightarrow$ 

DistributionOfComponentOfPhase

Creates a quantity representing the (molar) fraction of the specified component being present in the specified phase compared to the overall system [-]. This corresponds to the degree of segregation to that phase.

## Parameters

- phase The name of the phase
- **component** The name of the component

#### Returns

A new DistributionOfComponentOfPhase object.

## $\textbf{classmethod driving\_force\_for\_evaporation()} \rightarrow DrivingForceForEvaporation$

Creates a quantity representing the driving force for evaporation

## Returns

A new DrivingForceForEvaporation object.

## **classmethod dynamic\_viscosity**(*phase: str*) $\rightarrow$ DynamicViscosity

Creates a quantity representing the dynamic viscosity of a liquid phase.

## Parameters

**phase** – The name of the liquid phase

#### Returns

A new DynamicViscosity object.

classmethod electric\_conductivity(interface\_scattering\_constant=4e-08)  $\rightarrow$ ScheilElectricConductivity

Creates a quantity representing electric conductivity.

## Parameters

**interface\_scattering\_constant** – A constant describing the magnitude of the effect of phase interface scattering [ohm \* m]. Omitting it or giving a zero value means phase interface scattering is not considered.

#### Returns

A new ScheilElectricConductivity object.

**classmethod electric\_conductivity\_of\_phase**(*phase: str*)  $\rightarrow$  ElectricConductivityOfPhase

Creates a quantity representing the electric conductivity of a phase.

#### Parameters

phase – The name of the phase or ALL\_PHASES to choose all phases

#### Returns

A new ElectricConductivityOfPhase object.

**classmethod electric\_resistivity**(*interface\_scattering\_constant=4e-08*)  $\rightarrow$  ScheilElectricResistivity

Creates a quantity representing electric resistivity.

## Parameters

**interface\_scattering\_constant** – A constant describing the magnitude of the effect of phase interface scattering [ohm \* m]. Omitting it or giving a zero value means phase interface scattering is not considered.

## Returns

A new ScheilElectricResistivity object.

**classmethod electric\_resistivity\_of\_phase**(*phase: str*)  $\rightarrow$  ElectricResistivityOfPhase

Creates a quantity representing the electric resistivity of a phase.

## Parameters

phase – The name of the phase or ALL\_PHASES to choose all phases

## Returns

A new ElectricResistivityOfPhase object.

## $\texttt{classmethod evaporation\_enthalpy()} \rightarrow EvaporationEnthalpy$

Creates a quantity representing the evaporation enthalpy.

## Returns

A new EvaporationEnthalpy object.

## $\texttt{classmethod heat\_per\_gram()} \rightarrow HeatPerGram$

Creates a quantity representing the total heat release from the liquidus temperature down to the current temperature [J/g].

**Note:** The total or apparent heat release during the solidification process consists of two parts: one is the so-called latent heat, i.e. heat due to the liquid -> solid phase transformation (*latent\_heat\_per\_mole(*) and *latent\_heat\_per\_gram(*)), and the other is the heat related to the specific heat of liquid and solid phases (*heat\_per\_mole(*) and *heat\_per\_gram(*)).

#### Returns

A new HeatPerGram object.

## $\texttt{classmethod heat\_per\_mole()} \rightarrow HeatPerMole$

Creates a quantity representing the total heat release from the liquidus temperature down to the current temperature [J/mol].

**Note:** The total or apparent heat release during the solidification process consists of two parts: one is the so-called latent heat, i.e. heat due to the liquid -> solid phase transformation (*latent\_heat\_per\_mole(*) and *latent\_heat\_per\_gram(*)), and the other is the heat related to the specific heat of liquid and solid phases (*heat\_per\_mole(*) and *heat\_per\_gram(*)).

#### Returns

A new HeatPerMole object.

## **classmethod kinematic\_viscosity**(*phase: str*) $\rightarrow$ KinematicViscosity

Creates a quantity representing the kinematic viscosity of a liquid phase.

Parameters phase – The name of the liquid phase

Returns

A new KinematicViscosity object.

## $\texttt{classmethod latent\_heat\_per\_gram()} \rightarrow LatentHeatPerGram$

Creates a quantity representing the cumulated latent heat release from the liquidus temperature down to the current temperature [J/g].

**Note:** The total or apparent heat release during the solidification process consists of two parts: one is the so-called latent heat, i.e. heat due to the liquid -> solid phase transformation (*latent\_heat\_per\_mole(*) and *latent\_heat\_per\_gram(*)), and the other is the heat related to the specific heat of liquid and solid phases (*heat\_per\_mole(*) and *heat\_per\_gram(*)).

## Returns

A new LatentHeatPerGram object.

## $\texttt{classmethod latent\_heat\_per\_mole()} \rightarrow LatentHeatPerMole$

Creates a quantity representing the cumulated latent heat release from the liquidus temperature down to the current temperature [J/mol].

**Note:** The total or apparent heat release during the solidification process consists of two parts: one is the so-called latent heat, i.e. heat due to the liquid -> solid phase transformation (*latent\_heat\_per\_mole(*))

and latent\_heat\_per\_gram()), and the other is the heat related to the specific heat of liquid and solid phases (heat\_per\_mole() and heat\_per\_gram()).

#### Returns

A new LatentHeatPerMole object.

## **classmethod mass\_fraction\_of\_a\_solid\_phase**(*phase: str*) $\rightarrow$ MassFractionOfASolidPhase

Creates a quantity representing the mass fraction of a solid phase.

#### Parameters

**phase** – The name of the phase or ALL\_PHASES to choose all solid phases

#### Returns

A new MassFractionOfASolidPhase object.

#### $classmethod mass_fraction_of_all_liquid() \rightarrow MassFractionOfAllLiquid$

Creates a quantity representing the total mass fraction of all the liquid phase.

#### Returns

A new MassFractionOfAllLiquid object.

 $\texttt{classmethod mass\_fraction\_of\_all\_solid\_phases()} \rightarrow \texttt{MassFractionOfAllSolidPhase}$ 

Creates a quantity representing the total mass fraction of all solid phases.

#### Returns

A new MassFractionOfAllSolidPhase object.

#### classmethod molar\_mass\_of\_gas() $\rightarrow$ MolarMassOfGas

Creates a quantity representing the molar mass of gas

## Returns

A new MolarMassOfGas object.

#### **classmethod molar\_volume\_of\_phase**(*phase: str*) $\rightarrow$ MolarVolumeOfPhase

Creates a quantity representing the molar volume of a phase [m^3/mol].

#### Parameters

phase – The name of the phase or ALL\_PHASES to choose all phases

#### Returns

A new MolarVolumeOfPhase object.

#### $classmethod \ molar_volume_of_system() \rightarrow MolarVolumeOfSystem$

Creates a quantity representing the molar volume of the system [m<sup>3</sup>/mol].

#### Returns

A new MolarVolumeOfSystem object.

## **classmethod mole\_fraction\_of\_a\_solid\_phase**(*phase: str*) $\rightarrow$ MoleFractionOfASolidPhase

Creates a quantity representing the molar fraction of a solid phase.

## Parameters

phase – The name of the phase or ALL\_PHASES to choose all solid phases

#### Returns

A new MoleFractionOfASolidPhase object.

 $\texttt{classmethod mole_fraction_of_all_liquid()} \rightarrow \texttt{MoleFractionOfAllLiquid}$ 

Creates a quantity representing the total molar fraction of all the liquid phase.

Returns

A new MoleFractionOfAllLiquid object.

 $\texttt{classmethod mole_fraction_of_all_solid_phases()} \rightarrow \texttt{MoleFractionOfAllSolidPhases}$ 

Creates a quantity representing the total molar fraction of all solid phases.

#### Returns

A new MoleFractionOfAllSolidPhases object.

classmethod site\_fraction\_of\_component\_in\_phase(phase: str, component: str,

 $sub\_lattice\_ordinal\_no: int = 0) \rightarrow$ SiteFractionOfComponentInPhase

Creates a quantity representing the site fractions [-].

## **Parameters**

- phase The name of the phase, use ALL\_PHASES to choose all stable phases
- component The name of the component, use ALL\_COMPONENTS to choose all components
- **sub\_lattice\_ordinal\_no** The ordinal number (i.e. 1, 2, ...) of the sublattice of interest, use None to choose all sublattices

**Note:** Detailed information about the sublattices can be obtained by getting the *Phase* object of a phase from the *System* object using tc\_python.system.System.get\_phase\_in\_system. For each phase the sublattices are obtained by using tc\_python.system.Phase.get\_sublattices. The order in the returned list is equivalent to the sublattice ordinal number expected, **but note that the ordinal numbers start with 1**.

#### Returns

A new SiteFractionOfComponentInPhase object.

classmethod surface\_tension(*phase: str*)  $\rightarrow$  SurfaceTension

Creates a quantity representing the surface tension of a liquid phase.

#### Parameters

phase – The name of the liquid phase

#### Returns

A new SurfaceTension object.

classmethod temperature()  $\rightarrow$  Temperature

Creates a quantity representing the temperature [K].

#### Returns

A new Temperature object.

classmethod thermal\_conductivity(interface\_scattering\_constant=4e-08)  $\rightarrow$ 

ScheilThermalConductivity

Creates a quantity representing thermal conductivity.

#### Parameters

interface\_scattering\_constant – A constant describing the magnitude of the effect of

phase interface scattering [ohm \* m]. Omitting it or giving a zero value means phase interface scattering is not considered.

#### Returns

A new ScheilThermalConductivity object.

```
classmethod thermal_conductivity_of_phase(phase: str) \rightarrow ThermalConductivityOfPhase
```

Creates a quantity representing the thermal conductivity of a phase.

## Parameters

phase – The name of the phase or ALL\_PHASES to choose all phases

#### Returns

A new ThermalConductivityOfPhase object.

 $\texttt{classmethod thermal_diffusivity} (\textit{interface\_scattering\_constant=4e-08}) \rightarrow \texttt{ScheilThermalDiffusivity} (\textit{interface\_scattering\_scatteri$ 

Creates a quantity representing thermal diffusivity.

## Parameters

**interface\_scattering\_constant** – A constant describing the magnitude of the effect of phase interface scattering [ohm \* m]. Omitting it or giving a zero value means phase interface scattering is not considered.

#### Returns

A new ScheilThermalDiffusivity object.

## $\texttt{classmethod thermal_diffusivity_of_phase}(\textit{phase: str}) \rightarrow \texttt{ThermalDiffusivityOfPhase}$

Creates a quantity representing the thermal diffusivity of a phase.

## Parameters

phase – The name of the phase or ALL\_PHASES to choose all phases

#### Returns

A new ThermalDiffusivityOfPhase object.

**classmethod thermal\_resistivity**(*interface\_scattering\_constant=4e-08*)  $\rightarrow$  ScheilThermalResistivity Creates a quantity representing thermal resistivity.

#### **Parameters**

**interface\_scattering\_constant** – A constant describing the magnitude of the effect of phase interface scattering [ohm \* m]. Omitting it or giving a zero value means phase interface scattering is not considered.

#### Returns

A new ScheilThermalResistivity object.

## **classmethod thermal\_resistivity\_of\_phase**(*phase: str*) $\rightarrow$ ThermalResistivityOfPhase

Creates a quantity representing the thermal resistivity of a phase.

## Parameters

phase – The name of the phase or ALL\_PHASES to choose all phases

#### Returns

A new ThermalResistivityOfPhase object.

 $\texttt{classmethod volume\_fraction\_of\_a\_solid\_phase(\textit{phase: str}) \rightarrow VolumeFractionOfASolidPhase(phase: str) \rightarrow Volume$ 

Creates a quantity representing the volume fraction of a solid phase.

## **Parameters**

phase – The name of the phase or ALL\_PHASES to choose all solid phases

Returns

A new VolumeFractionOfASolidPhase object.

 $classmethod \ volume\_fraction\_of\_all\_liquid() \rightarrow VolumeFractionOfAllLiquid$ 

Creates a quantity representing the total volume fraction of all the liquid phase.

#### Returns

A new VolumeFractionOfAllLiquid object.

 $\texttt{classmethod volume\_fraction\_of\_all\_solid\_phases()} \rightarrow \texttt{VolumeFractionOfAllSolidPhases}()$ 

Creates a quantity representing the total volume fraction of all solid phases.

## Returns

A new VolumeFractionOfAllSolidPhases object.

#### class tc\_python.quantity\_factory.ThermodynamicQuantity

## Bases: AbstractQuantity

Factory class providing quantities used for defining equilibrium calculations (single equilibrium, property and phase diagrams, ...) and their results.

**Note:** In this factory class only the most common quantities are defined, you can always use the *Console Mode* syntax strings in the respective methods as an alternative (for example: "NPM(\*)").

**classmethod** activity\_of\_component(*component: str*, *use\_ser: bool* = *False*)  $\rightarrow$  ActivityOfComponent

Creates a quantity representing the activity of a component [-].

## Parameters

- component The name of the component, use ALL\_COMPONENTS to choose all components
- **use\_ser** Use Stable-Element-Reference(SER). The user-defined reference state is used if this setting is set to *False*.

#### Returns

A new ActivityOfComponent object.

**classmethod bulk\_modulus**(*phase: str*) → BulkModulus

Creates a quantity representing the bulk modulus of a phase.

#### Parameters

phase – The name of the phase or ALL\_PHASES to choose all phases

#### Returns

A new BulkModulus object.

classmethod chemical\_diffusion\_coefficient(phase: str, diffusing\_element: str, gradient\_element: str, reference\_element: str)  $\rightarrow$ 

## ChemicalDiffusionCoefficient

Creates a quantity representing the chemical diffusion coefficient of a phase [m^2/s].

## Parameters

- phase The name of the phase
- diffusing\_element The diffusing element
- gradient\_element The gradient element
- reference\_element The reference element (for example "Fe" in a steel)

## Returns

A new ChemicalDiffusionCoefficient object.

```
classmethod chemical_potential_of_component(component: str, use_ser: bool = False) \rightarrow ChemicalPotentialOfComponent
```

Creates a quantity representing the chemical potential of a component [J].

## Parameters

- component The name of the component, use ALL\_COMPONENTS to choose all components
- **use\_ser** Use Stable-Element-Reference(SER). The user-defined reference state is used if this setting is set to *False*.

#### Returns

A new ChemicalPotentialOfComponent object.

Creates a quantity representing the composition of a phase [mole-fraction].

#### **Parameters**

- phase The name of the phase, use ALL\_PHASES to choose all stable phases
- component The name of the component, use ALL\_COMPONENTS to choose all components

#### Returns

A new CompositionOfPhaseAsMoleFraction object.

```
\texttt{classmethod composition_of_phase_as_weight_fraction(\textit{phase: str, component: str})} \rightarrow
```

CompositionOfPhaseAsWeightFraction

Creates a quantity representing the composition of a phase [weight-fraction].

## **Parameters**

- phase The name of the phase, use ALL\_PHASES to choose all stable phases
- component The name of the component, use ALL\_COMPONENTS to choose all components

## Returns

A new CompositionOfPhaseAsWeightFraction object.

## **classmethod dynamic\_viscosity**(*phase: str*) $\rightarrow$ DynamicViscosity

Creates a quantity representing the dynamic viscosity of a liquid phase.

## Parameters

**phase** – The name of the liquid phase

## Returns

A new DynamicViscosity object.

## $\texttt{classmethod electric\_conductivity}() \rightarrow ElectricConductivity}$

Creates a quantity representing electric conductivity.

## Returns

A new ElectricConductivity object.

**classmethod electric\_conductivity\_of\_phase**(*phase: str*)  $\rightarrow$  ElectricConductivityOfPhase

Creates a quantity representing the electric conductivity of a phase.

#### Parameters

**phase** – The name of the phase or *ALL\_PHASES* to choose all phases

#### Returns

A new ElectricConductivityOfPhase object.

## classmethod electric\_resistivity() $\rightarrow$ ElectricResistivity

Creates a quantity representing electric resistivity.

#### Returns

A new ElectricResistivity object.

```
classmethod electric_resistivity_of_phase(phase: str) \rightarrow ElectricResistivityOfPhase
```

Creates a quantity representing the electric resistivity of a phase.

## Parameters

phase – The name of the phase or ALL\_PHASES to choose all phases

#### Returns

A new ElectricResistivityOfPhase object.

**classmethod gibbs\_energy\_of\_a\_phase**(*phase: str, use\_ser: bool* = *False*)  $\rightarrow$  GibbsEnergyOfAPhase Creates a quantity representing the Gibbs energy of a phase [J].

#### **Parameters**

- phase The name of the phase or ALL\_PHASES to choose all phases
- **use\_ser** Use Stable-Element-Reference(SER). The user-defined reference state will be used when this setting is set to False.

#### Returns

A new GibbsEnergyOfAPhase object.

**classmethod kinematic\_viscosity**(*phase: str*)  $\rightarrow$  KinematicViscosity

Creates a quantity representing the kinematic viscosity of a liquid phase.

#### **Parameters**

phase – The name of the liquid phase

#### Returns

A new KinematicViscosity object.

## classmethod mass\_fraction\_of\_a\_component(component: str) $\rightarrow$ MassFractionOfAComponent

Creates a quantity representing the mass fraction of a component.

#### Parameters

component - The name of the component or ALL\_COMPONENTS to choose all components

#### Returns

A new MassFractionOfAComponent object.

**classmethod mass\_fraction\_of\_a\_phase**(*phase: str*)  $\rightarrow$  MassFractionOfAPhase

Creates a quantity representing the mass fraction of a phase.

#### Parameters

**phase** – The name of the phase or *ALL\_PHASES* to choose all phases.

#### Returns

A new MassFractionOfAPhase object.

 $\texttt{classmethod molar_volume_of_phase}(\textit{phase: str}) \rightarrow MolarVolumeOfPhase}$ 

Creates a quantity representing the molar volume of a phase [m^3/mol].

## **Parameters**

phase – The name of the phase or ALL\_PHASES to choose all phases

#### Returns

A new MolarVolumeOfPhase object.

## $\texttt{classmethod molar_volume_of_system()} \rightarrow MolarVolumeOfSystem$

Creates a quantity representing the molar volume of the system [m^3/mol].

### Returns

A new MolarVolumeOfSystem object.

classmethod mole\_fraction\_of\_a\_component(component: str)  $\rightarrow$  MoleFractionOfAComponent

Creates a quantity representing the mole fraction of a component.

## Parameters

component – The name of the component or ALL\_COMPONENTS to choose all components

#### Returns

A new MoleFractionOfAComponent object.

classmethod mole\_fraction\_of\_a\_phase(*phase: str*)  $\rightarrow$  MoleFractionOfAPhase

Creates a quantity representing the mole fraction of a phase.

## Parameters

**phase** – The name of the phase or *ALL\_PHASES* to choose all phases

#### Returns

A new MoleFractionOfAPhase object.

#### classmethod normalized\_driving\_force\_of\_a\_phase(phase: str) $\rightarrow$

NormalizedDrivingForceOfAPhase

Creates a quantity representing normalized driving force of a phase [-].

**Warning:** A driving force calculation requires that the respective phase has been set to the state *DORMANT*. The parameter *All* is only reasonable if all phases have been set to that state.

#### **Parameters**

**phase** – The name of the phase or *ALL\_PHASES* to choose all phases

## Returns

A new DrivingForceOfAPhase object.

#### classmethod pressure() $\rightarrow$ Pressure

Creates a quantity representing the pressure [Pa].

#### Returns

A new Pressure object.

## classmethod shear\_modulus(phase: str) $\rightarrow$ ShearModulus

Creates a quantity representing the shear modulus of a phase.

#### **Parameters**

**phase** – The name of the phase or *ALL\_PHASES* to choose all phases

Returns A new ShearModulus object. classmethod surface\_tension(*phase: str*)  $\rightarrow$  SurfaceTension Creates a quantity representing the surface tension of a liquid phase. **Parameters phase** – The name of the liquid phase Returns A new SurfaceTension object. classmethod system\_size()  $\rightarrow$  SystemSize Creates a quantity representing the system size [mol]. Returns A new SystemSize object. classmethod temperature()  $\rightarrow$  Temperature Creates a quantity representing the temperature [K]. Returns A new Temperature object. **classmethod thermal\_conductivity()** → ThermalConductivity Creates a quantity representing thermal conductivity. Returns A new ThermalConductivity object. **classmethod** thermal\_conductivity\_of\_phase(*phase*: *str*)  $\rightarrow$  ThermalConductivityOfPhase Creates a quantity representing the thermal conductivity of a phase. **Parameters** phase – The name of the phase or ALL\_PHASES to choose all phases Returns A new ThermalConductivityOfPhase object. **classmethod thermal\_diffusivity**()  $\rightarrow$  ThermalDiffusivity Creates a quantity representing thermal diffusivity. Returns A new ThermalDiffusivity object. **classmethod thermal\_diffusivity\_of\_phase**(*phase: str*)  $\rightarrow$  ThermalDiffusivityOfPhase Creates a quantity representing the thermal diffusivity of a phase. **Parameters** 

**phase** – The name of the phase or *ALL\_PHASES* to choose all phases

#### Returns

A new ThermalDiffusivityOfPhase object.

classmethod thermal\_resistivity()  $\rightarrow$  ThermalResistivity

Creates a quantity representing thermal resistivity.

#### Returns

A new ThermalResistivity object.

**classmethod thermal\_resistivity\_of\_phase**(*phase: str*)  $\rightarrow$  ThermalResistivityOfPhase Creates a quantity representing the thermal resistivity of a phase.

**Parameters** 

phase – The name of the phase or ALL\_PHASES to choose all phases

Returns

A new ThermalResistivityOfPhase object.

**classmethod tracer\_diffusion\_coefficient**(*phase: str, diffusing\_element: str*)  $\rightarrow$ TracerDiffusionCoefficient

Creates a quantity representing tracer diffusion coefficient of a phase [m^2/s].

## Parameters

• phase – The name of the phase

• diffusing\_element - The diffusing element

#### Returns

A new TracerDiffusionCoefficient object.

classmethod u\_fraction\_of\_a\_component(component: str)  $\rightarrow$  UFractionOfAComponent

Creates a quantity representing the u-fraction of a component.

#### Parameters

component – The name of the component

#### Returns

A new UFractionOfAComponent object.

#### classmethod user\_defined\_function(*expression: str*) $\rightarrow$ Function

Creates a quantity representing a user-defined function.

Parameters expression – The function expression

#### Returns

A new Function object

**classmethod volume\_fraction\_of\_a\_phase**(*phase: str*)  $\rightarrow$  VolumeFractionOfAPhase

Creates a quantity representing the volume fraction of a phase.

#### Parameters

phase – The name of the phase or ALL\_PHASES to choose all phases

## Returns

A new VolumeFractionOfAPhase object.

## $\texttt{classmethod youngs\_modulus}(\mathit{phase: str}) \rightarrow YoungsModulus}$

Creates a quantity representing the Young's modulus of a phase.

## Parameters

phase - The name of the phase or ALL\_PHASES to choose all phases

## Returns

A new YoungsModulus object.

## 6.6 Module "utils"

```
class tc_python.utils.CompositionType(value)
     Bases: Enum
     The type of composition.
     COMPONENT = 1
          Composition given per component, this will be identical to ELEMENT in case of metals.
     ELEMENT = 0
          Composition given per element.
class tc_python.utils.CompositionUnit(value)
     Bases: Enum
     The composition unit.
     MASS_FRACTION = 1
          Mass fraction.
     MASS_PERCENT = 0
          Mass percent.
     MOLE_FRACTION = 3
          Mole fraction.
     MOLE PERCENT = 2
          Mole percent.
class tc_python.utils.ConversionUnit(value)
     Bases: Enum
     The composition unit used in a conversion.
     MOLE_FRACTION = 0
          Mole fraction.
     MOLE_PERCENT = 1
          Mole percent.
     WEIGHT_FRACTION = 2
          Weight fraction.
     WEIGHT_PERCENT = 3
          Weight percent.
class tc_python.utils.GasAmountUnit(value)
     Bases: Enum
     The amount of a gas.
     KILOGRAM = 1
          Kilogram.
     NORM_CUBIC_METER = 0
          Norm cubic meter (according to ISO 2533, p=101325 Pa, T=288.15 K, typically used for trading of gas
          bottles). Other definitions vary only slightly.
```

```
class tc_python.utils.GasCompositionUnit(value)
```

Bases: Enum

The composition unit for a gas.

```
MASS_FRACTION = 1
```

Mass fraction.

 $MASS\_PERCENT = 0$ 

Mass percent.

```
MOLE\_FRACTION = 3
```

Mole fraction.

 $MOLE_PERCENT = 2$ 

Mole percent.

## VOLUME\_FRACTION = 5

Volume fraction.

## $VOLUME_PERCENT = 4$

Volume percent.

## class tc\_python.utils.GasRateUnit(value)

Bases: Enum

The rate of a gas flow.

## $KILOGRAM_PER_SEC = 1$

Kilogram per second.

## NORM\_CUBIC\_METER\_PER\_SEC = 0

Norm cubic meter per second (according to ISO 2533, p=101325 Pa, T=288.15 K, typically used for trading of gas bottles). Other definitions vary only slightly.

## class tc\_python.utils.InterfacePosition(value)

Bases: Enum

The position of an interface relative to its region. Only used for diffusion simulations.

## LOWER = 0

The interface is on the lower side of its region.

## UPPER = 1

The interface is on the upper side of its region.

## class tc\_python.utils.PhaseUnit(value)

Bases: Enum

The units available for a phase fraction.

## $MASS_FRACTION = 1$

Mass fraction.

## $MOLE\_FRACTION = 0$

Mole fraction.

## VOLUME\_FRACTION = 2

Volume fraction.

## class tc\_python.utils.ResultValueGroup(result\_line\_group\_java)

## Bases: object

A x-y-dataset representing a line data calculation result (i.e. a Thermo-Calc quantity 1 vs. quantity 2).

**Warning:** Depending on the calculator, the dataset might contain *NaN*-values to separate the data between different subsets.

## Variables

- label a str describing what the data corresponds to
- **x** list of floats representing the first quantity ("x-axis")
- y list of floats representing the second quantity ("y-axis")

## $get_label() \rightarrow str$

Accessor for the line label :return the line label

## $get_x() \rightarrow List[float]$

Accessor for the x-values :return the x values

 $\textbf{get\_y()} \rightarrow List[float]$ 

Accessor for the y-values :return the y values

## class tc\_python.utils.TemperatureProfile

Bases: object

Represents a time-temperature profile used by non-isothermal calculations.

**Note:** The total simulation time can differ from the defined temperature profile. Constant temperature is assumed for any timepoint after the end of the defined profile.

## add\_time\_temperature(time: float, temperature: float)

Adds a time-temperature point to the non-isothermal temperature profile.

## Parameters

- time The time [s]
- temperature The temperature [K]

## Returns

This TemperatureProfile object

## 6.7 Module "propertymodel\_sdk"

class tc\_python.propertymodel\_sdk.AddPrefixToWarning(name=")

Bases: Filter

## filter(record)

Determine if the specified record is to be logged.

Returns True if the record should be logged, or False otherwise. If deemed appropriate, the record may be modified in-place.

```
class tc_python.propertymodel_sdk.CCTResult(quantity_id: str, description: str)
```

Bases: ResultQuantity

Represents a Continuous Cooling (CCT) result.

## Parameters

- quantity\_id The id of this result
- description The description of this result

add\_time\_temperature(time\_temperature\_id: str, description: str)

Adds a time-temperature pair to the result.

## Parameters

- time\_temperature\_id The id of the time-temperature pair
- description The description of the time-temperature pair

## temperature\_suffix = ' (T)'

The temperature suffix of a CCTResult

```
time_suffix = ' (t)'
```

The time suffix of a *CCTResult* 

**class** tc\_python.propertymodel\_sdk.**CCTResultValues**(*cooling\_rate: float* = -1.0,

```
cooling_rate_start_temperature: float = -1.0,
cooling_rate_end_temperature: float = -1.0)
```

Bases: object

Represents Continuous Cooling (CCT) result values.

Parameters

- **cooling\_rate** The cooling rate [K/s]
- cooling\_rate\_start\_temperature The start temperature of cooling [K]
- cooling\_rate\_end\_temperature The end temperature of cooling [K]

set\_result\_time\_temperature(time\_temperature\_id: str, time: float, temperature: float)
Sets a time-temperature pair of the result.

## **Parameters**

- time\_temperature\_id The id of the time-temperature pair
- time The time [s]
- temperature The temperature [K]

class tc\_python.propertymodel\_sdk.CalculationContext(system: System, model\_utils=None)

## Bases: object

Represents the interface of the Property Model with the Thermo-Calc application and the rest of the TC-Python functionality.

## Parameters

- system The system object of this calculation
- model\_utils The model utils object

#### $get_argument_ids() \rightarrow Set[str]$

Returns a list with the arguments of the models. Including arguments created from dynamic parameters.

#### get\_dependent\_component() $\rightarrow$ str

Obtains the dependent component from the UI

Note: The dependent component is that which has no composition specified explicitly, typically this is the major element of the material (such as Fe, Al, Ni,  $\ldots$ )

### Returns

The dependent component

#### get\_element\_names\_in\_camel\_case()

Obtains the elements in the system. The dependent component is not included.

#### Returns

A list of the elements in the system, each written in camel case.

## **get\_mass\_fractions()** → Dict[str, float]

Obtains the current composition from the UI as mass-fraction.

**Note:** In case of stepping over one or multiple axis, the returned data will represent the composition at the current step.

#### Returns

The composition (key: component, value: content) [mass-fraction]

## $\texttt{get\_mass\_percents()} \rightarrow Dict[str, float]$

Obtains the current composition from the UI in mass-percent.

**Note:** In case of stepping over one or multiple axis, the returned data will represent the composition at the current step.

#### Returns

The composition (key: component, value: content) [mass-percent]

## $get_model_path() \rightarrow str$

Obtains the current path to the Property Models folder defined by the user.

#### Returns

The absolute path to the Property Models folder defined by the user.

## $\texttt{get\_mole\_fractions()} \rightarrow \texttt{Dict[str, float]}$

Obtains the current composition from the UI as mole-fraction.

**Note:** In case of stepping over one or multiple axis, the returned data will represent the composition at the current step.

#### Returns

The composition (key: component, value: content) [mole-fraction]

## $get_mole_percents() \rightarrow Dict[str, float]$

Obtains the current composition from the UI in mole-percent.

**Note:** In case of stepping over one or multiple axis, the returned data will represent the composition at the current step.

#### Returns

The composition (key: component, value: content) [mole-percent]

## get\_temperature() $\rightarrow$ float

Obtains the current temperature from the UI.

#### Returns

The temperature [K]

#### get\_ui\_boolean\_value(component\_id: str) → bool

Obtains the value from the specified checkbox UI component.

Parameters component\_id – Id of the checkbox

#### Returns

The setting of the checkbox

#### get\_ui\_condition\_list(component\_id: str) → ConditionListEntry

Used to get the selected condition from components of type UIConditionListComponent :param component\_id: Id of the list UI component :return: The selected condition

#### get\_ui\_float\_value(component\_id: str) → float

Obtains the value from the specified UI component.

#### Parameters

component\_id – Id of the UI component

## Returns

The value

#### **get\_ui\_list\_value**(*component\_id: str*) → str

Obtains the selected entry from a UI component list. If a special element (such as ANY, NONE, ...) is selected, the corresponding locale-independent placeholder is provided.

#### Parameters

 $\textbf{component\_id} - Id \ of \ the \ list \ UI \ component$ 

#### Returns

The selected entry

## get\_ui\_string\_value(component\_id: str) → str

Obtains the selected entry from a UI component text field.

#### Parameters

**component\_id** – Id of the string UI component

#### Returns

The selected entry

## 

Obtains the temperature from the specified temperature UI component.

## Parameters

component\_id – Id of the temperature UI component

#### Returns

The temperature [K], note that input unit of the UI is specified in the model panel. If required, the temperature is automatically converted to K.

set\_result\_cct\_values(quantity\_id: str, r: CCTResultValues)

Sets the value of a previously defined result quantity (of type *CCTResultValues*) for further usage in the Thermo-Calc application for plotting, etc.

#### **Parameters**

- quantity\_id unique id of the result quantity
- **r** the CCTResultValues to be set

set\_result\_quantity\_value(quantity\_id: str, value: float, parameter: str = ")

Sets the value of a previously defined result quantity for further usage in the Thermo-Calc application for plotting, etc.

**Note:** Any result quantity that remains unset is automatically set to *NaN*.

#### **Parameters**

- quantity\_id Unique id of the result quantity
- parameter Use if result is parameterized. f.i. "per phase"
- **value** The value to be set

set\_single\_equilibrium\_result(quantity\_id: str, r: SingleEquilibriumResult)

## class tc\_python.propertymodel\_sdk.ConditionListEntry

Bases: object

Used in combination with components of type UIConditionListComponent.

Contains the element, if the selected condition is a composition Contains the Console Mode syntax of the selected condition. Contains the unit of the selected condition

## class tc\_python.propertymodel\_sdk.PropertyModel(\_locale: str = 'en-US')

Bases: object

The abstract base class for all property models.

**Note:** Every Property Model needs to implement most of the abstract methods of this class. However, some abstract methods are optional and should only be implemented if required.

**Note:** If overwriting the constructor in a Property Model, the constructor of the implemented class must have the identical signature and should pass the parameters to this base class constructor.

**Tip:** It is possible to switch off **internal INFO-log messages coming from the calculation engine** by changing the log-level on the TC-Python log object like this: *logging.getLogger("tc\_python").setLevel(logging.ERROR)*.

## **Parameters**

\_locale – The locale to be used, this is an internal parameter and is of no meaning to the end-user

## Variables

**logger** – logger object that is connected to the Thermo-Calc UI (*INFO*- and *WARNING*-level will be printed as *INFO*, *ERROR*-level as *ERROR*), it can be accessed like this: *self.logger.info*("*Some message*")

**abstract** add\_button\_callback(component\_id: str, ui\_components: Dict[UIComponent, int]) → List[UIComponent]

Implement this method if you have one or more UI components on which you called UIComponent. enable\_add\_button(), which adds a + button next to the component.

This method will be executed when you press any such + button.

This method is typically used to add more UI components dynamically and the method must return a list of the UI components to be added.

## This method can optionally be implemented by a Property Model.

#### Parameters

- component\_id The id of the UI component next to the pressed + button
- **ui\_components** dict with the components and the index of the current ui components of the model. Including arguments created in previous calls to add\_button\_callback.

## Returns

A list of UIComponent objects to be added

## TODO: docstring

## abstract after\_evaluations()

Called by the Thermo-Calc application immediately after the last model evaluation (using the method *PropertyModel.evaluate\_model()*). Use this method for any required cleanup.

## This method can optionally be implemented by a Property Model.

## abstract before\_evaluations(context: CalculationContext)

Called by the Thermo-Calc application immediately before the first model evaluation (using the method *PropertyModel.evaluate\_model()*). Use this method for any required preparations.

#### This method can optionally be implemented by a Property Model.

#### **Parameters**

 $\textbf{context} - The \ calculation \ context$ 

#### abstract evaluate\_model(context: CalculationContext)

Called by the Thermo-Calc application when the model should be actually calculated. **This is the mainmethod of the Property Model that contains the actual calculation code.** 

This method needs to be implemented by all property models.

## Parameters

**context** – The calculation context, this provides access to the Thermo-Calc application and all other TC-Python modules

## **abstract get\_license\_key()** $\rightarrow$ str

Provides the license key of the model.

## This method can optionally be implemented by a Property Model.

## abstract provide\_calculation\_result\_quantities() $\rightarrow$ List[*ResultQuantity*]

Called by the Thermo-Calc application when the model should provide its result quantity objects.

## This method needs to be implemented by all property models.

## Returns

Result quantity objects of the model (to be filled later with results in the method *PropertyModel.evaluate\_model()*)

## abstract provide\_model\_category() $\rightarrow$ List[str]

Called by the Thermo-Calc application when the model should provide its category (shown in the Thermo-Calc model tree).

## This method needs to be implemented by all property models.

#### Returns

Category of the model, it may be present in several categories

## abstract provide\_model\_description() $\rightarrow$ str

Called by the Thermo-Calc application when the model should provide its detailed description.

## This method needs to be implemented by all property models.

## Returns

Description text for the model

## <code>abstract provide\_model\_name()</code> $\rightarrow$ str

Called by the Thermo-Calc application when the model should provide its name (shown in the Thermo-Calc model tree).

## This method needs to be implemented by all property models.

## Returns

Name of the model

## abstract provide\_model\_parameters() $\rightarrow$ Dict[str, float]

Called by the Thermo-Calc application when the model should provide all model parameters and their current values.

## This method can optionally be implemented by a Property Model.

**Note:** These are internal variables of the Property Model that are intended to be modified from the outside. Typically this is used to adjust their values in a optimizer during the development of the model.

## Returns

The model parameter ids and their current values [unit according to the parameter meaning]

## abstract provide\_ui\_panel\_components() → List[UIComponent]

Called by the Thermo-Calc application when the model should provide its UI components for the model panel to be plotted. This happens also whenever a model gets checked in the model tree.

## This method needs to be implemented by all property models.

## Returns

Model UI panel components in the order to be presented in the model panel

# **abstract remove\_button\_callback**(*component\_id: str, ui\_components: Dict*[UIComponent, *int*]) $\rightarrow$ List[str]

Implement this method if you have one or more UI components on which you called UIComponent. enable\_remove\_button(), which adds a - button next to the component.

This method will be executed when you press any such - button.

This method is typically used to remove UI components dynamically and the method must return a list of the ids of the components that are going to be removed.

## This method can optionally be implemented by a Property Model.

## Parameters

- **component\_id** the id of the UI component next to the pressed button
- **ui\_components** list with the components and the index of the current ui components of the model. Including arguments created in previous calls to add\_button\_callback.

#### Returns

a list of UI component ids that are required to be removed

## abstract set\_model\_parameter(model\_parameter\_id: str, value: float)

Called by the Thermo-Calc application when a model parameter should be reset.

## This method can optionally be implemented by a Property Model.

**Note:** These are internal variables of the Property Model that are intended to be modified from the outside. Typically this is used to adjust their values in a optimizer during the development of the model.

## **Parameters**

- model\_parameter\_id The parameter id
- **value** The value [unit according to the parameter meaning]

## Bases: object

Defines a calculation result quantity of a Property Model that is identified by a unique id.

## **Parameters**

- **quantity\_id** Unique id of the quantity
- description Description of the quantity (shown in the Thermo-Calc UI)
- **quantity\_type** Type of the quantity (defines the unit)

 $\begin{array}{c} \texttt{get\_description()} \rightarrow \texttt{str} \\ & \texttt{Obtains the description of the quantity.} \\ & \texttt{Returns} \\ & \texttt{Description of the quantity} \\ & \texttt{get\_id()} \rightarrow \texttt{str} \\ & \texttt{Obtains the id of the quantity.} \end{array}$ 

Returns Unique id of the quantity

**get\_type()**  $\rightarrow$  *ResultQuantityType* Obtains the type of quantity.

## Returns

Type of the quantity

## class tc\_python.propertymodel\_sdk.ResultQuantityType(value)

Bases: Enum

Defining the type of a result quantity.

CCT\_QUANTITY = 5

A cct quantity

#### $ENERGY_QUANTITY = 2$

An energy quantity

## GENERAL\_QUANTITY = 0

A general quantity

#### LENGTH\_QUANTITY = 7

A length in quantity

## SINGLE\_EQUILIBRIUM\_QUANTITY = 6

A cct quantity

## SOLIDIFICATION\_RATE\_QUANTITY = 12

A solidification rate quantity

## STRAIN\_QUANTITY = 14

A strain quantity

## STRENGTH\_QUANTITY = 8

A strength quantity

## STRESS\_QUANTITY = 13

A stress quantity

## SURFACE\_ENERGY\_QUANTITY = 3

A surface energy quantity

## **TEMPERATURE\_QUANTITY = 1**

A temperature quantity

## $THERMAL\_GRADIENT\_QUANTITY = 11$

A thermal gradient quantity
## $TIME_QUANTITY = 4$

A time quantity

## $VELOCITY_QUANTITY = 10$

A velocity quantity

## VICKERS\_HARDNESS\_QUANTITY = 9

A hardness quantity

## class tc\_python.propertymodel\_sdk.SpecialListMarkers

Bases: object

Placeholders for special list elements that are locale-dependent. They will be provided by UI list components if a special marker has been selected.

## ANY\_LIST\_MARKER = 'ANY'

Marker that represents "Any"

## NONE\_LIST\_MARKER = 'NONE'

Marker that represents "None"

## Bases: UIComponent

Checkbox UI component of the model panel.

## **Parameters**

- component\_id Unique id of the component
- name Name of the component, will be presented in the model panel
- description Additional description of the component
- **setting** Initial setting of the checkbox

## connect\_component\_enableability(dependent\_component\_id: str)

Connects the state enabled of any other UI component of the model panel to the value of this boolean component.

## **Parameters**

**dependent\_component\_id** – Id of the UI element to be dependent on this boolean component

## connect\_component\_visibility(dependent\_component\_id: str)

Connects the visibility of any other UI component of the model panel to the value of this boolean component.

## Parameters

**dependent\_component\_id** – Id of the UI element to be dependent on this boolean component

## enable\_add\_button()

Adds a + button to the right of the UI component.

## Returns

This UI component

## enable\_remove\_button()

Adds a - button to the right of the UI component.

## Returns

This UI component

## $\texttt{get\_dependent\_components()} \rightarrow List[str]$

Obtains a list containing all UI elements currently connected regarding their visibility.

## Returns

A list with the component id of all UI elements currently connected

## $\texttt{get\_setting()} \rightarrow \texttt{bool}$

Obtains the setting of the checkbox.

## Returns

The setting of the checkbox

## remove\_component\_visibility(dependent\_component\_id: str)

Removes the visibility connection to a UI component that has been previously connected.

Parameters dependent\_component\_id – Id of the previously connection UI element

## set\_index: int = -1)

Sets the position in the graphical user interface.

Parameters index – The position

## Returns

This UI component

## class tc\_python.propertymodel\_sdk.UIComponent(component\_id: str, name: str, description: str)

## Bases: object

Abstract Base class for all UI components of the model panel.

Never make an instance of UIComponent, always use the sub-classes. For instance UIStringComponent.

## **Parameters**

- component\_id Unique id of the component
- name Name of the component, will be presented in the model panel
- **description** Additional description of the component

## $\texttt{get\_description()} \rightarrow str$

Obtains the additional description of the component.

## Returns

Additional description of the component

## $\texttt{get\_id}() \to str$

Obtains the unique id of the component.

## Returns

Unique id of the component

#### $get_name() \rightarrow str$

Obtains the name of the component.

#### Returns

Name of the component, will be presented in the model panel

## set\_enabled(enabled)

Sets whether the component should be enabled or not

## Parameters enabled -

Returns This UI component

#### set\_has\_callback(has\_callback)

## set\_visible(visible)

Sets the visibility of the component

# Parameters visible –

Returns

This UI component

#### Bases: UIComponent

System condition list UI component of the model panel.

Parameters

- component\_id Unique id of the component
- name Name of the component, will be presented in the model panel
- description Additional description of the component

class tc\_python.propertymodel\_sdk.UIFloatComponent(component\_id: str, name: str, description: str,

value: float)

## Bases: UIComponent

General real value text field UI component of the model panel.

## Parameters

- component\_id Unique id of the component
- name Name of the component, will be presented in the model panel
- description Additional description of the component
- value Initial setting of the text field

## enable\_add\_button()

Adds a + button to the right of the UI component.

#### Returns

This UI component

## enable\_remove\_button()

Adds a - button to the right of the UI component.

## Returns

this UI component

 $\texttt{get\_value()} \to \texttt{float}$ 

Obtains the setting of the text field.

#### Returns

The setting of the text field

set\_index: int = -1)

Sets the position in the graphical user interface.

#### Parameters index – The position

Returns

This UI component

## Bases: UIComponent

General list UI component of the model panel that can contain any strings.

#### **Parameters**

- component\_id Unique id of the component
- name Name of the component, will be presented in the model panel
- description Additional description of the component
- **content** Entries of the list, they need to contain a locale-independent id and a localized content string, for example: [("ENTRY\_1\_ID", "entry 1"), (ENTRY\_2\_ID", "entry 2")]
- **selected\_entry** Entry to be initially selected. If omitted, by default the first element is selected.

**connect\_component\_visibility**(dependent\_component\_id: str, selected\_item\_to\_set\_visible: str)

Connects the visibility of any other UI component of the model panel to the selection of a certain entry of the list.

#### **Parameters**

- **dependent\_component\_id** Id of the UI element to be dependent on the chosen element
- **selected\_item\_to\_set\_visible** Entry (locale independent id) of the list to be chosen to set the dependent component visible

## enable\_add\_button()

Adds a + button to the right of the UI component.

#### Returns

This UI component

## enable\_remove\_button()

Adds a - button to the right of the UI component.

## Returns

This UI component

## $get\_content() \rightarrow List[Tuple[str, str]]$

Obtains the entries of the list.

## Returns

Entries of the list, they need to contain a locale-independent id and a localized content string, for example: [("ENTRY\_1\_ID", "entry 1"), (ENTRY\_2\_ID", "entry 2")]

## $\texttt{get\_dependent\_components()} \rightarrow Dict[str, List[str]]$

Obtains a dictionary containing all UI elements currently connected regarding their visibility.

## Returns

All UI elements currently connected (key: dependent component id, value: required list entries to set it visible)

## get\_selected\_entry() $\rightarrow$ str

Obtains the initially selected entry.

## Returns

Initially selected entry. If empty, the first element is selected.

## remove\_component\_visibility(dependent\_component\_id: str)

Removes the visibility connection to a UI component that has been previously connected.

#### Parameters

dependent\_component\_id - Id of the previously connection UI element

#### set\_index(index: int = -1)

Sets the position in the graphical user interface.

Parameters index – The position

## Returns

This UI component

## Bases: UIComponent

Phase list UI component of the model panel.

#### Parameters

- **component\_id** Unique id of the component
- name Name of the component, will be presented in the model panel
- description Additional description of the component
- **default\_phase** Default phase, if omitted no default phase is chosen and only initially the first element of the list is selected. If an ANY-marker is added, this is chosen as the default element.
- **any\_marker\_setting** Defines if an entry "ANY PHASE" should be added to the phase list, if set to true this overrides any default phase setting

#### enable\_add\_button()

Adds a + button to the right of the UI component.

## Returns

This UI component

#### enable\_remove\_button()

Adds a - button to the right of the UI component.

#### Returns

This UI component

## $\texttt{get\_any\_marker\_setting()} \rightarrow \texttt{bool}$

Obtains the setting if any entry "ANY PHASE" is added to the phase list.

#### Returns

If an entry "ANY PHASE" is added to the phase list, if set to true this overrides any default phase setting

## $\texttt{get\_default\_phase()} \rightarrow str$

Obtains the default phase.

## Returns

Default phase, if omitted no default phase is chosen and only initially the first element of the list is selected. If an ANY-marker is added, this is chosen as the default element.

#### set\_index(index: int = -1)

Sets the position in the graphical user interface.

Parameters index – The position

#### Returns

This UI component

## class tc\_python.propertymodel\_sdk.UISectionDividerComponent(component\_id: str, name: str,

description: str)

## Bases: UIComponent

General text field UI component of the model panel.

## Parameters

- component\_id Unique id of the component
- name Name of the component, will be presented in the model panel
- description Additional description of the component
- string Initial setting of the text field

## $\texttt{get\_value()} \rightarrow \textit{str}$

Obtains the setting of the text field.

#### Returns

The setting of the text field

## set\_index(index: int = -1)

Sets the position in the graphical user interface.

## Parameters

**index** – The position

#### Returns

This UI component

#### Bases: UIComponent

General text field UI component of the model panel.

#### **Parameters**

- component\_id Unique id of the component
- name Name of the component, will be presented in the model panel
- description Additional description of the component
- string Initial setting of the text field

## enable\_add\_button()

Adds a + button to the right of the UI component.

## Returns

This UI component

## enable\_remove\_button()

Adds a - button to the right of the UI component.

#### Returns

This UI component

## $get_value() \rightarrow str$

Obtains the setting of the text field.

#### Returns

The setting of the text field

#### set\_index(index: int = -1)

Sets the position in the graphical user interface.

#### **Parameters**

index - The position

#### Returns

This UI component

## Bases: UIComponent

Temperature value text field UI component of the model panel.

#### Parameters

- component\_id Unique id of the component
- name Name of the component, will be presented in the model panel
- description Additional description of the component
- **temp** Initial temperature to be set in the text field (unit defined by the user in the Thermo-Calc system)

## enable\_add\_button()

Adds a + button to the right of the UI component.

## Returns

This UI component

## enable\_remove\_button()

Adds a - button to the right of the UI component.

## Returns

This UI component

## $\texttt{get\_temp()} \to \texttt{float}$

Obtains the temperature set in the text field.

## Returns

The temperature to be set in the text field (unit defined by the user in the Thermo-Calc system)

## **set\_index**(*index*: *int* = -1)

Sets the position in the graphical user interface.

Parameters index – The position

## Returns

This UI component

 $\label{eq:component_id:str,name:str,description:str,initial_setting:bool} tc_python.propertymodel_sdk.create_boolean_ui_component(component_id:str,name:str,description:str,initial_setting:bool) \rightarrow UIBooleanComponent$ 

Creates a UI checkbox component for a boolean value. The value of that component can later be accessed during the model evaluation.

## Parameters

- component\_id Unique id of the component
- name Name of the component, will be presented in the model panel
- description Additional description of the component
- initial\_setting Initial setting of the checkbox

## Returns

The created component

tc\_python.propertymodel\_sdk.create\_condition\_list\_ui\_component(component\_id: str, name: str, description: str)  $\rightarrow$ 

**UIConditionListComponent** 

Creates a UI list component for all conditions defined in the system. The value of that component can later be accessed during the model evaluation.

## Parameters

- component\_id Unique id of the component
- **name** Name of the component, will be presented in the model panel
- description Additional description of the component

## Returns

The created component

tc\_python.propertymodel\_sdk.create\_energy\_quantity(quantity\_id: str, description: str)  $\rightarrow$ ResultQuantity

Creates a UI energy result quantity (in J). When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

## Parameters

- quantity\_id Unique id of the result quantity
- description Additional description of the result quantity

## Returns

The created result quantity

tc\_python.propertymodel\_sdk.create\_float\_ui\_component(component\_id: str, name: str, description:  $str, value: float) \rightarrow UIFloatComponent$ 

Creates a UI text field component for a real number. The value of that component can later be accessed during the model evaluation.

#### Parameters

- component\_id Unique id of the component
- name Name of the component, will be presented in the model panel
- description Additional description of the component
- **value** Initial setting of the text field

## Returns

The created component

tc\_python.propertymodel\_sdk.create\_general\_quantity(quantity\_id: str, description: str) →

#### **ResultQuantity**

Creates a general result quantity that can contain any type of result (without a unit). When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

## Parameters

- quantity\_id Unique id of the result quantity
- description Additional description of the result quantity

## Returns

The created result quantity

```
tc_python.propertymodel_sdk.create_length_quantity(quantity_id: str, description: str) \rightarrow
ResultQuantity
```

Creates a length result quantity. When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

## Parameters

- **quantity\_id** Unique id of the result quantity
- description Additional description of the result quantity

## Returns

The created result quantity

tc\_python.propertymodel\_sdk.create\_list\_ui\_component(component\_id: str, name: str, description: str, entry\_list: List[Tuple[str, str]], selected\_entry:  $str = ") \rightarrow UIGeneralListComponent$  Creates a UI list component for string entries. The value of that component can later be accessed during the model evaluation.

## Parameters

- component\_id Unique id of the component
- name Name of the component, will be presented in the model panel
- description Additional description of the component
- **entry\_list** Entries of the list, they need to contain a locale-independent id and a localized content string, for example: [("ENTRY\_1\_ID", "entry 1"), (ENTRY\_2\_ID", "entry 2")]
- **selected\_entry** Entry to be initially selected. If omitted, by default the first element is selected.

## Returns

The created component

Creates a UI list component for all phases defined in the system. It is possible to select a default phase that is supposed to be the **expected phase selection** for that list. The value of that component can later be accessed during the model evaluation.

A **default** phase is the phase that is initially selected and re-selected as soon as a currently selected phase is removed. If the default phase is not available, a "NONE"-marker will be created and used instead of the default phase. A typical use case for the default phase setting is a phase list that expects to contain the LIQUID-phase of a system.

## Parameters

- component\_id Unique id of the component
- name Name of the component, will be presented in the model panel
- description Additional description of the component
- **default\_phase** Default phase, if omitted no default phase is chosen and only initially the first element of the list is selected. **If an ANY-marker is added, this is chosen as the default element.**
- **any\_marker** Defines if an entry "ANY PHASE" should be added to the phase list, if set to true this overrides any default phase setting

## Returns

The created component

tc\_python.propertymodel\_sdk.create\_section\_divider\_ui\_component(component\_id: str, name: str, description: str)  $\rightarrow$ UISectionDividerComponent

Creates an empty UI component acting as a spacer. :param component\_id: Unique id of the component :return: The created component

tc\_python.propertymodel\_sdk.create\_solidification\_rate\_quantity(quantity\_id: str, description: str)  $\rightarrow ResultQuantity$ 

Creates a Solidification rate result quantity. When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

## Parameters

- quantity\_id Unique id of the result quantity
- description Additional description of the result quantity

## Returns

The created result quantity

```
tc_python.propertymodel_sdk.create_strain_quantity(quantity_id: str, description: str) \rightarrow
ResultQuantity
```

Creates a strain result quantity. When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

## **Parameters**

- quantity\_id Unique id of the result quantity
- description Additional description of the result quantity

## Returns

The created result quantity

tc\_python.propertymodel\_sdk.create\_strength\_quantity(quantity\_id: str, description: str)  $\rightarrow$ ResultQuantity

Creates a strength result quantity. When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

## **Parameters**

- **quantity\_id** Unique id of the result quantity
- description Additional description of the result quantity

## Returns

The created result quantity

 $\texttt{tc_python.propertymodel\_sdk.create\_stress\_quantity(quantity\_id: str, description: str) \rightarrow \texttt{tc_python.propertymodel\_sdk.create\_stress\_quantity(quantity\_id: str, description: str))} \rightarrow \texttt{tc_python.propertymodel\_sdk.create\_stress\_quantity(quantity\_id: str, description: str))}$ 

## *ResultQuantity*

Creates a stress result quantity. When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

## Parameters

- **quantity\_id** Unique id of the result quantity
- description Additional description of the result quantity

## Returns

The created result quantity

tc\_python.propertymodel\_sdk.create\_string\_ui\_component(component\_id: str, name: str, description: str, string: str)  $\rightarrow$  UIStringComponent

Creates a UI text field component. The value of that component can later be accessed during the model evaluation.

## **Parameters**

- **component\_id** Unique id of the component
- name Name of the component, will be presented in the model panel
- description Additional description of the component
- **string** Initial setting of the text field

## Returns

The created component

```
tc_python.propertymodel_sdk.create_surface_energy_quantity(quantity_id: str, description: str) \rightarrow ResultQuantity
```

Creates an energy result quantity (in J). When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

## **Parameters**

- **quantity\_id** Unique id of the result quantity
- description Additional description of the result quantity

## Returns

The created result quantity

```
tc_python.propertymodel_sdk.create_temperature_quantity(quantity_id: str, description: str) \rightarrow
ResultOuantity
```

Creates a temperature result quantity (in K). When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

## **Parameters**

- **quantity\_id** Unique id of the result quantity
- description Additional description of the result quantity

## Returns

The created result quantity

Creates a UI text field component for a temperature value. The value of that component can later be accessed during the model evaluation.

## Parameters

- **component\_id** Unique id of the component
- **name** Name of the component, will be presented in the model panel
- description Additional description of the component
- **initial\_temp** Initial temperature to be set in the text field. (The unit of initial\_temp is Kelvin. The value in the text field will be automatically converted using the unit chosen by the user.)

## Returns

The created component

 $\texttt{tc_python.propertymodel\_sdk.create\_thermal\_gradient\_quantity(quantity\_id: str, description: str) \rightarrow ResultQuantity}$ 

Creates a Thermal gradient result quantity. When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

## **Parameters**

- **quantity\_id** Unique id of the result quantity
- description Additional description of the result quantity

## Returns

The created result quantity

```
tc_python.propertymodel_sdk.create_time_quantity(quantity_id: str, description: str) \rightarrow ResultQuantity
```

Creates a time result quantity (in s). When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

## **Parameters**

- quantity\_id Unique id of the result quantity
- description Additional description of the result quantity

## Returns

The created result quantity

```
tc_python.propertymodel_sdk.create_velocity_quantity(quantity_id: str, description: str) \rightarrow
ResultQuantity
```

Creates a Velocity result quantity. When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

## Parameters

- quantity\_id Unique id of the result quantity
- description Additional description of the result quantity

## Returns

The created result quantity

 $\texttt{tc_python.propertymodel_sdk.create_vickers_hardness_quantity}(\textit{quantity\_id: str, description: str}) \rightarrow \textit{ResultQuantity}$ 

Creates a Vickers hardness result quantity. When the model is evaluated, a value can be added to the quantity and it will be used to transfer the result to the Thermo-Calc plot engine.

## Parameters

- **quantity\_id** Unique id of the result quantity
- description Additional description of the result quantity

## Returns

The created result quantity

## 6.8 Module "exceptions"

## exception tc\_python.exceptions.APIServerException

Bases: GeneralException

An exception that occurred during the communication with the API-server. It is normally not related to an error in the user program.

## exception tc\_python.exceptions.CalculationException

Bases: TCException

An exception that occurred during a calculation.

## exception tc\_python.exceptions.ComponentNotExistingException

Bases: GeneralException

The selected component is not existing.

#### exception tc\_python.exceptions.DatabaseException

Bases: CalculationException

Error loading a thermodynamic or kinetic database, typically due to a misspelled database name or a database missing in the system.

#### exception tc\_python.exceptions.DegreesOfFreedomNotZeroException

Bases: CalculationException

The degrees of freedom in the system are not zero, i.e. not all required conditions have been defined. Please check the conditions given in the exception message.

## exception tc\_python.exceptions.EquilibriumException

Bases: CalculationException

An equilibrium calculation has failed, this might happen due to inappropriate conditions or a very difficult problem that can not be solved.

#### exception tc\_python.exceptions.GeneralCalculationException

Bases: CalculationException

General error occurring while a calculation is performed.

## exception tc\_python.exceptions.GeneralException

Bases: TCException

A general exception that might occur in different situations.

#### exception tc\_python.exceptions.InvalidCalculationConfigurationException

Bases: CalculationException

Thrown when errors are detected in the configuration of the calculation.

#### exception tc\_python.exceptions.InvalidCalculationStateException

Bases: CalculationException

Trying to access an invalid calculation object that was invalidated by calling *invalidate* on it.

#### exception tc\_python.exceptions.InvalidNumberOfResultGroupsException

Bases: ResultException

A calculation result contains several result groups, which is not supported for the used method.

## exception tc\_python.exceptions.InvalidResultConfigurationException

Bases: ResultException

A calculation result configuration is invalid.

## exception tc\_python.exceptions.InvalidResultStateException

Bases: CalculationException

Trying to access an invalid result (for example a SingleEquilibriumTempResult object that got already invalidated by condition changes or a result that was invalidated by calling *invalidate* on it).

## exception tc\_python.exceptions.LicenseException

Bases: GeneralException

No valid license for the API or any Thermo-Calc product used by it found.

## exception tc\_python.exceptions.NoDataForPhaseException

Bases: ResultException

There is no result data available for a selected phase.

## exception tc\_python.exceptions.NotAllowedOperationException

Bases: CalculationException

The called method or operation is not allowed in the current mode of operation (i.e. debug or production mode). *Production mode* means that the Property Model is only present as an \*.*py.encrypted*-file, while in *debug mode* it is available as \*.*py*-file. Certain methods for obtaining internal model parameters are not available for encrypted models.

#### exception tc\_python.exceptions.PhaseNotExistingException

Bases: GeneralException

The selected phase is not existing, so no data can be provided for it.

#### exception tc\_python.exceptions.ResultException

Bases: TCException

An exception that occurred during the configuration of a calculation result.

## exception tc\_python.exceptions.SyntaxException

Bases: CalculationException

Syntax error in a Console Mode expression.

## exception tc\_python.exceptions.TCException

Bases: Exception

The root exception of TC-Python.

## exception tc\_python.exceptions.UnrecoverableCalculationException

Bases: CalculationException

The calculation reached a state where no further actions are possible, this happens most often due to a FORTRANhard crash in the API server backend.

**Note:** It is possible to catch that exception outside of the *with*-clause context and to continue by setting up a new context (i.e. by a new *with TCPython() as session*).

tc\_python.exceptions.handle\_exception(e)

## 6.9 Module "abstract\_base"

## class tc\_python.abstract\_base.AbstractCalculation(calculator)

Bases: object

Abstract base class for calculations.

## get\_configuration\_as\_string() $\rightarrow$ str

Returns detailed information about the current state of the calculation object.

**Warning:** The structure of the calculator objects is an implementation detail and might change between releases without notice. **Therefore do not rely on the internal object structure**.

#### get\_system\_data() $\rightarrow$ SystemData

Returns the content of the database for the currently loaded system. This can be used to modify the parameters and functions and to change the current system by using *with\_system\_modifications()*.

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as \*.tdb-file.

#### Returns

The system data

## invalidate()

Invalidates the object and frees the disk space used by it. *This is only required if the disk space occupied by the object needs to be released during the calculation*. No data can be retrieved from the object afterwards.

## with\_system\_modifications(system\_modifications: SystemModifications)

Updates the system of this calculator with the supplied system modification (containing new phase parameters and system functions).

**Note:** This is only possible if the system has been read from unencrypted (i.e. *user*) databases loaded as a \*.tdb-file.

#### Parameters

system\_modifications – The system modification to be performed

#### Returns

## class tc\_python.abstract\_base.AbstractResult(result)

Bases: object

Abstract base class for results. This can be used to query for specific values .

## invalidate()

Invalidates the object and frees the disk space used by it. *This is only required if the disk space occupied by the object needs to be released during the calculation*. No data can be retrieved from the object afterwards.

## class tc\_python.abstract\_base.PhaseParameter(parameter\_name: Union[str, object])

## Bases: object

Database phase parameter expression used by SystemModifications.set().

## Parameters

parameter\_name - The phase parameter name

#### get\_intervals() → List[TemperatureInterval]

Returns the list of all defined intervals.

#### Returns

The defined temperature intervals

#### get\_lower\_temperature\_limit() $\rightarrow$ float

Returns the lower temperature limit.

#### Returns

The lower temperature limit in K

## $get_name() \rightarrow str$

Returns the name of the phase parameter.

#### Returns

The name of the phase parameter.

## remove\_all\_intervals()

Removes all previously defined temperature intervals.

#### Returns

This PhaseParameter object

#### remove\_interval\_with\_upper\_limit(upper\_temperature\_limit: float)

Removes a previously defined temperature interval with matching upper temperature limit.

If no such interval exists, an exception is thrown.

#### Returns

This PhaseParameter object

#### 

Adds/overwrites a parameter expression for a temperature interval.

## **Default value of the upper limit of the interval**: 6000 K

**Note:** The lower temperature limit is either defined by the lower temperature limit given with *PhaseParameter.set\_lower\_temperature\_limit()* or by the upper temperature limit of the adjacent interval.

**Note:** If there is an existing interval with exactly the same *upper\_temperature\_limit*, that interval is over-written, otherwise the interval is added.

## **Parameters**

- **parameter\_expression** The parameter expression, example: +V34\*T\*LN(T)+V35\*T\*\*2+V36\*T\*\*(-1)+V37\*T\*\*3")
- **upper\_temperature\_limit** The upper temperature limit for which the expression should be used

#### Returns

This PhaseParameter object

#### set\_interval(interval: TemperatureInterval)

Adds/overwrites a temperature interval.

**Note:** The lower temperature limit is either defined by the lower temperature limit given with *PhaseParameter.set\_lower\_temperature\_limit()* or by the upper temperature limit of the adjacent

interval.

**Note:** If there is an existing interval with exactly the same *upper\_temperature\_limit*, that interval is over-written, otherwise the interval is added.

#### Returns

This PhaseParameter object

#### set\_lower\_temperature\_limit(lower\_temperature\_limit: float = 298.15)

Sets the lower temperature limit of the phase parameter.

Default: 298.15 K

Parameters lower\_temperature\_limit – The lower temperature limit in K

#### Returns

This *PhaseParameter* object

## class tc\_python.abstract\_base.SystemData(system\_data)

Bases: object

Provides information about the parameters and functions of a user database. The obtained objects can be used to modify the database using with\_system\_modifications() of all calculators.

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as \*.tdb-file.

## get\_phase\_parameter(parameter: str) $\rightarrow$ PhaseParameter

Returns a phase parameter.

Example:

system\_data.get\_phase\_parameter('G(HCP\_A3,FE:VA;0)')

Note: Parameters can only be read from unencrypted (i.e. user) databases loaded as a \*.tdb-file.

**Note:** For details about the syntax search the Thermo-Calc help for *GES* (the name for the Gibbs Energy System module in Console Mode).

#### Parameters

**parameter** – The name of the phase parameter (for example: "*G*(*LIQUID*,*FE*;0)")

#### Returns

The phase parameter

## $\texttt{get\_phase\_parameter\_names()} \rightarrow List[str]$

Returns all phase parameters present in the current system.

#### Returns

The list of phase parameters

## $get_system_function(f: str) \rightarrow SystemFunction$

Returns a system function.

Note: The parameter 'f' was previously called 'function' but was renamed.

Example:

system\_data.get\_system\_function('GHSERCR')

Note: Functions can only be read from unencrypted (i.e. user) databases loaded as a \*.tdb-file.

**Note:** For details about the syntax search the Thermo-Calc help for *GES* (the name for the Gibbs Energy System module in Console Mode).

## **Parameters**

**f** – The name of the system function (for example: "GHSERCR")

#### Returns

The system function

## get\_system\_function\_names() $\rightarrow$ List[str]

Returns all system functions present in the current system.

#### Returns

The list of system functions

## class tc\_python.abstract\_base.SystemFunction(function\_name: Union[str, object])

## Bases: object

Database function expression used by SystemModifications.set().

#### Parameters

function\_name – The function name

## $get_intervals() \rightarrow List[TemperatureInterval]$

Returns the list of all defined intervals.

#### Returns

The defined temperature intervals

#### get\_lower\_temperature\_limit() $\rightarrow$ float

Returns the lower temperature limit.

## Returns

The lower temperature limit in K

## $\texttt{get\_name()} \to \mathsf{str}$

Returns the name of the system function.

## Returns

The name of the system function

#### remove\_all\_intervals()

Removes all previously defined temperature intervals.

#### Returns

This SystemFunction object

## remove\_interval\_with\_upper\_limit(upper\_temperature\_limit: float)

Removes a previously defined temperature interval with matching upper temperature limit.

#### If no such interval exists, an exception is thrown.

#### Returns

This SystemFunction object

**set\_expression\_with\_upper\_limit**(*function\_expression: str, upper\_temperature\_limit: float* = 6000.0) Adds/overwrites a function expression for a temperature interval.

#### Default value of the upper limit of the interval: 6000 K

**Note:** The lower temperature limit is either defined by the lower temperature limit given with *SystemFunction.set\_lower\_temperature\_limit()* or by the upper temperature limit of the adjacent interval.

**Note:** If there is an existing interval with exactly the same *upper\_temperature\_limit*, that interval is over-written, otherwise the interval is added.

#### Parameters

- function\_expression The function expression, example: +V34\*T\*LN(T)+V35\*T\*\*2+V36\*T\*\*(-1)+V37\*T\*\*3")
- **upper\_temperature\_limit** The upper temperature limit for which the expression should be used

#### Returns

This SystemFunction object

#### set\_interval(interval: TemperatureInterval)

Adds/overwrites a temperature interval.

**Note:** The lower temperature limit is either defined by the lower temperature limit given with *SystemFunction.set\_lower\_temperature\_limit()* or by the upper temperature limit of the adjacent interval.

**Note:** If there is an existing interval with exactly the same *upper\_temperature\_limit*, that interval is over-written, otherwise the interval is added.

#### Returns

This SystemFunction object

set\_lower\_temperature\_limit(lower\_temperature\_limit: float = 298.15)

Sets the lower temperature limit of the system function.

**Default**: 298.15 K

## Parameters

lower\_temperature\_limit - The lower limit in K

Returns

This SystemFunction object

## class tc\_python.abstract\_base.SystemModifications

Bases: object

Functionality to modify a user database during a calculation by changing phase parameters and system functions.

The actual changes are **only applied** by using tc\_python.abstract\_base.AbstractCalculation. with\_system\_modifications() on a calculator object.

## run\_ges\_command(ges\_command: str)

Sends a GES-command. This is actually applied when running `with\_system\_modifications` on a calculator object.

Example:  $run_ges\_command("AM-PH-DE FCC\_A1 C\_S 2 Fe:C")$  for adding a second composition set to the FCC\_A1 phase with Fe as major constituent on first sublattice and C as major constituent on second sublattice.

**Note:** For details about the syntax search the Thermo-Calc help for *GES* (the name for the Gibbs Energy System module in Console Mode).

**Note:** It should not be necessary for most users to use this method, try to use the corresponding method implemented in the API instead.

**Warning:** As this method runs raw GES-commands directly in the engine, it may hang the program in case of spelling mistakes (e.g. forgotten parenthesis, ...).

## Parameters

**ges\_command** – The GES-command (for example: "AM-PH-DE FCC\_A1 C\_S 2 Fe:C")

## Returns

This SystemModifications object

set(parameter\_or\_function: Union[PhaseParameter, SystemFunction])

Overwrites or creates a phase parameter or system function.

Example: system\_modifications.set(PhaseParameter('G(LIQUID,FE;0)').set\_expression\_with\_upper\_limit('+1.2\*GFELIQ

Example: system\_modifications.set(SystemFunction("DGDEF").set\_expression\_with\_upper\_limit(+10.0-R\*T', 1000).set\_expression\_with\_upper\_limit(+20.0-R\*T', 3000))

Note: The old parameter/function is overwritten and any temperature intervals not defined are lost.

Note: Please consult the Thermo-Calc GES-system documentation for details about the syntax.

#### Returns

This SystemModifications object

class tc\_python.abstract\_base.TemperatureInterval(expression: Union[str, object],

upper\_temperature\_limit: float)

## Bases: object

Temperature interval expression used by the classes *SystemFunction* and *PhaseParameter*.

## **Parameters**

- expression The temperature function expressed in Thermo-Calc database syntax.
- upper\_temperature\_limit The upper temperature limit in K

## $\texttt{get\_expression()} \rightarrow str$

Returns the function expression of this temperature interval.

Returns

The temperature function expression

## $\texttt{get\_upper\_temperature\_limit()} \to \texttt{float}$

Returns the upper limit of this temperature interval.

## Returns

The upper temperature limit in K

## set\_expression(expression: str)

Sets the function expression of this temperature interval.

## Parameters expression – The temperature function expression

## set\_upper\_temperature\_limit(upper\_temperature\_limit: float)

Sets the upper limit of this temperature interval.

## Parameters

upper\_temperature\_limit - The upper temperature limit in K

## 6.10 Module "license"

## class tc\_python.license.LicenseManager(license\_manager)

Bases: object

Manages the license operations for Thermo-calc licenses for TC-Python, including login, activation, and proxy settings.

## activate(user: str, password: str)

Activates the license.

## Parameters

- **user** The username
- password The password

## activate\_offline(user: str, password: str, path: str = ")

Activate the license with an offline activation response file.

## **Parameters**

- **user** The username
- **password** The password
- path Optional path to folder where the activation response file is located

**create\_offline\_activation\_file**(*user: str, password: str, path: str = ''*)

Create a file for offline activation.

## Parameters

- **user** The username
- password The password
- path Optional path to folder where the activation file is created

## deactivate()

Deactivates the license if the license was activated using online method.

## Raises

**LicenseException** – If not logged in or license was activated using activate\_offline().

#### deactivate\_offline(path: str = ")

Deactivates the license if the license was activated using *activate\_offline(*).

#### Parameters

path – Optional path to folder where the deactivation file is located

## Raises

LicenseException - If not logged in

## $\texttt{get\_info()} \to \mathsf{str}$

Retrieves the current license information.

#### Returns

The license information

#### proxy\_settings\_remove()

Disables the proxy settings.

## proxy\_settings\_set(host: str, port: int, user: str, password: str)

Sets the proxy settings with the given host, port, username, and password.

## **Parameters**

- host The proxy host
- port The proxy port
- user The proxy username
- password The proxy password

## update()

Updates the license.

## CHAPTER

## SEVEN

## TROUBLESHOOTING

This section provides an FAQ for common problems that occur when using TC-Python.

## 7.1 Diagnostics script

If you have problems running TC-Python, run the diagnostics script below.

On Linux you can alternatively download the script directly into your current working directory by:

```
.....
```

```
Run this script when troubleshooting TC-Python
It is important to run this script EXACTLY the same way as you run your TC-Python script
(In the same IDE, same project, same Python environment, same Jupyter notebook e.t.c)
.....
version = '2025b'
print('Testing TC-Python version: ' + version)
print('Please make sure that the variable "version" above, matches the release that you.
↔want to test, if not change it and re-run this script.')
# below this line, nothing needs to be manually updated.
import sys
print('')
print('Python version (needs to be at least Python 3.8, Python 2.x is not supported):')
print(sys.version)
if sys.version_info[0] < 3 or sys.version_info[1] < 8:</pre>
    print('Wrong version of Python !!!!!')
print('')
print('Python executable path: (gives a hint about the used virtual / conda environment,
\rightarrow in case of Anaconda the corresponding n'
      'environment name can be found by running `conda env list` on the Anaconda command.
→prompt, '
                                                                             (continues on next page)
```

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```
'TC-Python must be installed into \nEACH separate environment used!)')
print(sys.executable)
import os
print('')
print('Thermo-Calc ' + version + ' installation directory: (must be a valid path to a.
Gomplete installation of ' + version + ')')
tc_env_variable = 'TC' + version[2:].upper() + '_HOME'
try:
   print(os.environ[tc_env_variable])
except:
   print('No Thermo-calc environment variable for ' + version + ' was found. (' + tc_

env_variable + ')')

import tc_python
numerical_version = version[:-1]
if version[-1] == 'a':
   numerical_version += '.1.*'
elif version[-1] == 'b':
   numerical_version += '.2.*'
print('')
print('TC-Python version: (needs to be ' + numerical_version + ')')
print(tc_python.__version__)
user_based_license_var = os.environ.get('TC_LICENSE_SPRING', None)
user_based_license = False
if user_based_license_var is not None:
   user_based_license = user_based_license_var.upper() == 'Y'
if not user_based_license:
   print('Url of license server: (if license server is NO-NET, you need a local license.
→file)')
   trv:
       print(os.environ['LSHOST'])
   except:
        print('No Thermo-calc license server url was found. (LSHOST)')
   print('')
   print('Path to local license file: (only necessary if not using license server)')
   try:
       print(os.environ['LSERVRC'])
   except:
        print('No path to local license file was found. (LSERVRC)')
else:
   print('')
   print('User/password based licenses is enabled')
   print("License Information:")
   with tc_python.TCPython() as session:
        license_manager = session.get_license_manager()
        print(license_manager.get_info())
```

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```
with tc_python.TCPython() as session:
    print('')
    print('Lists the databases (should be a complete list of the installed databases_
    othat you have license for or do not require license):')
    print(session.get_databases())
```

## 7.2 "No module named tc\_python" error on first usage

This problem occurs because your used Python interpreter cannot find the TC-Python package. We expect that you have installed the TC-Python package in your **Python system interpreter** following the instructions in the *Installation Guide*.

Normally the error message "*No module named tc\_python*" is caused by unintentionally configuring a PyCharm project to use a so-called **Virtual Environment**. This happens unfortunately by default when creating a new PyCharm project with not changing the default settings.

**Note:** A Virtual Environment is basically a separate and completely independent copy of the system-wide Python interpreter. It does not contain any packages.

On Windows systems we recommend to use the Anaconda Python Distribution as Python interpreter. However, the instructions given here are valid for any operating system and distribution.

Since TC-Python 2018b we do recommend to **not use Virtual Environments** unless there is a reasonable use case for that.

There are two possible solutions to fix the problem:

1. The quick fix for your problem is to run

pip install <path to the TC-Python folder>/TC\_Python-<version>-py3-none-any.whl

within the *Terminal window* of the opened PyCharm project. This *Terminal window* automatically runs within the *Virtual Environment* configured for the project (if any). You can see the name of the *Virtual Environment* at the beginning of each command prompt line (here it is called *venv*):

Microsoft Windows [Version 10.0.16299.431] (c) 2017 Microsoft Corporation. All rights reserved.

```
(venv) C:\Users\User\Documents\>
```

The command will consequently **install TC-Python also within that Virtual Environment automatically**. The Terminal window can be found at the bottom of the IDE. Note that it might be necessary to enable these buttons first by selecting the menu entry **View** $\rightarrow$ **Tool Buttons**.

- 2. The better fix is to change your project **to use the system interpreter**. This is described in detail in the section *Fixing potential issues with the environment* in Step 5 of the *Installation Guide*.
  - It is recommendable to use that approach also for all your future projects.

Both fixes will only change the configuration of the opened project. Further useful information can be found in the section *Python Virtual Environments*.

## 7.3 "pip install" fails with "Failed to establish a new network connection" or similar

If *pip install* fails with a network related error (might also be "*socket not available*", "*retrying after connection broken*", …) it is often due to the computer being behind a proxy-server, this is common in large organizations. Of course also the network connection might be broken.

TC-Python has dependencies to a few other packages:

- *py4j*
- jproperties
- *six* (transient dependency of *jproperties*)
- 1. The recommended approach is to simply use *pip*. It will resolve the dependencies automatically by downloading them from the *PyPI*-repository server (https://pypi.org). If your computer is located behind a proxy-server, the connection to the repository will fail. In that case it is necessary to configure *pip* with the detailed configuration of the proxy server:

pip install -proxy user:password@proxy\_ip:port py4j jproperties

2. Another alternative is to manually download the latest \*.whl-file of each dependency from the repository server (https://pypi.org -> *Search projects*) and to install it manually using:

```
pip install py4j-#.#.#-py2.py3-none-any.whl
...
```

The actual actual version number needs to be inserted into the file name. The downside of this approach is that updates to that package have to be fully manual also in the future. Additionally it is also necessary to install all transient dependencies in that way.

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