TC-Python Documentation

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Thermo-Calc Software AB

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CHAPTER

ONE

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.

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 3.9.18 interpreter is containing the following major packages:

Package	Version
matplotlib	3.8
numpy	1.26.0
scikit-learn	1.3.1
scipy	1.11.3
pyvista	0.42.2
TC-Python	2024a

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 *Mac OS: 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\2024a\python\python.exe
Linux	/home/UserName/Thermo-Calc/2024a/python/bin/python3
MacOS	/Applications/Thermo-Calc-2024a.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\2024a\SDK\
		TC-Python
Linux		/home/UserName/Thermo-Calc/2024a/SDK/TC-Python
MacOS		/Users/Shared/Thermo-Calc/2024a/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, \dots), 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 tem	sys-	Path to the TC-Python folder
Windows		C:\Users\UserName\Documents\Thermo-Calc\2024a\SDK\ TC-Python
Linux		/home/UserName/Thermo-Calc/2024a/SDK/TC-Python
MacOS		/Users/Shared/Thermo-Calc/2024a/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 *Mac OS: 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\2024a\SDK\
		TC-Python
Linux		/home/UserName/Thermo-Calc/2024a/SDK/TC-Python
MacOS		/Users/Shared/Thermo-Calc/2024a/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 constall c
```

However, that should normally not be required.

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

pip show TC-Python

CHAPTER

TWO

MAC OS: SETTING ENVIRONMENT VARIABLES

In order to use TC-Python on Mac you need to set some environment variables.

TC24A_HOME=/Applications/Thermo-Calc-2024a.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

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.

CHAPTER

THREE

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.

3.1 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')
```

3.2 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.

3.3 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.

3.3.1 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")
            )
```

3.3.2 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)
```

3.3.3 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:

3.3.4 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.

3.3.5 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"))
        )
```

3.3.6 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:

3.3.7 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)
```

3.3.8 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")
```

3.3.9 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
```

3.3.10 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=0)
    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'] }")
```

3.3.11 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*.

3.4 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")
```

3.4.1 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.

```
# ...
# some code to set up a Diffusion calculation
# ...
first_diffusion_result = diffusion_calculation.calculate()
continued_calculation = first_diffusion_result.with_continued_calculation()
continued_calculation.set_simulation_time(110000.0)
continued_calculation.with_left_boundary_condition(BoundaryCondition.mixed_zero_flux_and_
--activity().set_activity_for_element('C', 1.0))
second_result = continued_calculation.calculate()
# ...
# Now you can use get second_result to get calculated values, just as normal.
# You can also use first_diffusion_result even after second_result is created.
# You can also use second_result (and even first_diffusion_result) to create a new_
--ContinuedDiffusionCalculation by calling with_continued_calculation.
```

3.5 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.

3.5.1 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.

3.5.2 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).

3.5.3 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\2024a\PropertyModels
Linux	/home/UserName/Thermo-Calc/2024a/PropertyModels or: /opt/Thermo-Calc/
	2024a/PropertyModels
MacOS	<pre>/Applications/Thermo-Calc-2024a.app/Contents/Resources/PropertyModels</pre>

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

BEST PRACTICES

4.1 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()))
```

4.2 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)
```

4.3 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 ...
```

4.4 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.

4.5 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)
```

4.6 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().
```

```
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
```

4.7 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:
```

```
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...')
```

4.8 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		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)	nt Logout			
File Edit	View Insert Cell Kernel Widgets Help	Python [conda env:Anaconda3] O			
₽ + %	42 € ♦ UN E C > Code				
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>				
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>				

4.9 Property Model Framework

4.9.1 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		
14				normatized_nitving_iorce)	
75	ging mode	ι:			
76 c_python.implementationpropertymodel_sdk import _PropertyModelProxy					
77	<pre>77 rtyModelProxy(model=DrivingForcePythonModel(), java_port=57334, python_port=65247)</pre>				
		Ŭ			
				🗹 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:

- 🖧 DrivingForcePythonModel.py 🗵			
pe 1 from tc_python import *			
3			
4 <mark> </mark>	PropertyModel):		
5 RESULT_QUANTITY = "driving			
6 NORMALIZED_RESULT_QUANTITY	= "normalizedDrivingForce		
7 MATRIX_PHASE_LIST = "matrix			
8 PRECIPITATE_PHASE_LIST = "	precipitate"		
9	Show Context Actions	Alt+Enter	
10 🔿 🖯 def provide_calculation_re	D <u>P</u> aste	Ctrl+V	ntity]:
11 🖕 results = [create_ener	Copy / Paste Special		ULT_QUANTITY,
12	Column Selection <u>M</u> ode	Alt+Shift+Insert	uantity.text")),
13 create_gene	Find <u>U</u> sages	Alt+F7	RMALIZED_RESULT_QUANTITY,
14 🛱 🔤	<u>R</u> efactor		<pre>izedResultQuantity.text"))]</pre>
15 A return results	Folding		
16	Go To	•	
17 of def provide_model_category	Generate	Alt+Insert	
18 A return [selfget_prop	Run 'DrivingForcePythonMo'	Ctrl+Shift+F10	
19	Debug 'DrivingForcePythonMo'	Curtonnetrio	
20 🔿 🔁 def provide_model_name(sel	-	Coverage	
	🕻 Run 'DrivingForcePythonMo' with	r 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()
....
```

4.9.2 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.

4.9.2.1 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:
```

__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.

4.9.2.2 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**.

4.9.3 Alternative Python for Property Models

4.9.3.1 Default bundled Python interpreter

Thermo-Calc is by default using a Python 3.9.18 interpreter bundled to the software for running the property models. It is containing the following major packages:

Package	Version
matplotlib	3.8
numpy	1.26.0
scikit-learn	1.3.1
scipy	1.11.3
pyvista	0.42.2
TC-Python	2024a

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	:\Program Files\Thermo-Calc\2024a\python\python.exe				
Linux	/home/UserName/Thermo-Calc/2024a/python/bin/python3				
MacOS	/Applications/Thermo-Calc-2024a.app/Contents/Resources/python/				
	bin/python3				

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

4.10 Process Metallurgy Calculations

4.10.1 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 ...
```

4.10.2 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.

4.10.3 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.

4.10.4 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

FIVE

API REFERENCE

5.1 Calculations

5.1.1 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

$get_components() \rightarrow List[str]$

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).

5.1.2 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

$\texttt{get_components()} \rightarrow List[str]$

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_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.

5.1.3 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]

A value that adds to the activation energy of mobility data from the database. Default: 0.0 J/mol

Parameters

mobility_enhancement_activation_energy – The value that adds to the activation energy of mobility data from the database [J/mol].

set_mobility_enhancement_prefactor(*mobility_enhancement_prefactor*: *float* = 1.0)

A parameter that multiplies to the mobility data from database. Default: 1.0

Parameters mobility_enhancement_prefactor – The mobility enhancement factor [-]

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 m³/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]

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_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.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.

Returns

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

Returns

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: 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 – The interfacial energy [J/m^2]

Returns This PrecipitatePhase object

Note: The calculation of the interfacial energy using a broken-bond model is based on the assumption of an interface between a bcc- and a fcc-crystal structure with (110) and (111) lattice planes regardless of the actual phases.

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⁻³].

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⁻³].

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 m⁴/(Js).

Parameters

phase_boundary_mobility – The phase boundary mobility [m⁴/(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

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*.

 $\texttt{get_aspect_ratio_distribution_for_particle_length_of(\textit{precipitate_id: str, time: float}) \rightarrow \texttt{float}) \rightarrow \texttt{float} \rightarrow \texttt{$

[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

time.

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

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

Returns

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

$\texttt{get_cubic_factor_distribution_for_radius_of(\textit{precipitate_id: str, time: float}) \rightarrow [\texttt{List[float]}, \texttt{List[float]}) \rightarrow [\texttt{List[float]}) \rightarrow [\texttt{List[float]}, \texttt{List[float]}) \rightarrow [\texttt{List[float]}, \texttt{List[float]}) \rightarrow [\texttt{List[float]}, \texttt{List[float]}) \rightarrow [\texttt{List[float]}, \texttt{List[float]}) \rightarrow [\texttt{List[float]}) \rightarrow [\texttt{List[float]}, \texttt{List[float]}) \rightarrow [\texttt{List[float]}) \rightarrow [\texttt{Li$

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)

get_grain_critical_radius() \rightarrow [List[float], 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])

get_grain_mean_radius() → [List[float], 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]

Returns

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

$get_grain_size_distribution(\mathit{time: float}) \rightarrow [List[float], List[float]]$

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)

get_matrix_composition_in_weight_fraction_of(*element_name: str*) \rightarrow [List[float], 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)

get_mean_particle_length_of(precipitate_id: str) → [List[float], 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

Returns

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])

$\texttt{get_normalized_grain_size_distribution(\textit{time: float}) \rightarrow [\texttt{List[float]},\texttt{List[float]}]$

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)

 $\texttt{get_normalized_number_density_distribution_of(precipitate_id: str, time: float) \rightarrow [\texttt{List[float]}, \texttt{list[float]}) \rightarrow [\texttt{List[float]}) \rightarrow [\texttt{List[float]}, \texttt{list[float]}) \rightarrow [\texttt{List[float]}) \rightarrow [\texttt{List[float]}, \texttt{list[float]}) \rightarrow [\texttt{List[float]}, \texttt{list[float]}) \rightarrow [\texttt{List[float]}, \texttt{list[float]}) \rightarrow [\texttt{List[float]}, \texttt{list[float]}) \rightarrow [\texttt{List[float]}) \rightarrow$

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 $[m^{-2}]$)

```
\label{eq:get_number_density_distribution_2d_for_radius_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 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])

get_number_density_distribution_for_particle_length_of(precipitate_id: str, time: float) → [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

Returns

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 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

Returns

A tuple of two lists of floats (radius [m], number of particles per unit area per unit length $[m^{-3}]$)

 $\label{eq:get_size_distribution_for_particle_length_of(\textit{precipitate_id: str, time: float}) \rightarrow [List[float]], \\ List[float]]$

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 $[m^{-4}]$)

 $\texttt{get_size_distribution_for_radius_of}(\textit{precipitate_id: str, time: float}) \rightarrow [\texttt{List[float]}, \texttt{List[float]}]$

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_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 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)

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 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 UnrecoverableCalculationException 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]

Returns

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(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%).

5.1.4 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)

Returns

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.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.

Returns

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, and ScheilSoluteTrapping enables solute trapping in the solid primary phase. :param scheil_type: Type of Scheil calculation, either ScheilClassic, ScheilBackDiffusion or ScheilSoluteTrapping :return: 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

$\texttt{get_stable_phases()} \rightarrow \texttt{List[str]}$

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) \rightarrow 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

$get_values_grouped_by_stable_phases_of(x_quantity: Union[ScheilQuantity, str], y_quantity: Union[ScheilQuantity, str], sort_and_merge: bool = True)$ $<math>\rightarrow$ 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. :return: A *ScheilBackDiffusion*

classmethod scheil_classic()

Configuration for Classic Scheil with fast diffusers. :return: 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. :return: 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.

Returns

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. 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_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

5.1.5 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

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

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

$get_components() \rightarrow List[str]$

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.

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

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

$\texttt{get_components()} \rightarrow List[str]$

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

Returns

The component names

$\texttt{get_gibbs_energy_addition_for}(\textit{phase: str}) \rightarrow \texttt{float}$

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

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 Returns

Parameters

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	
max_no_of_iterations – 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*')

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

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)

$get_tie_lines() \rightarrow ResultValueGroup$

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

$\texttt{get_x()} \rightarrow \texttt{List[float]}$

Accessor for the x-value :return: the x value

```
get_y() \rightarrow List[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

5.1.6 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.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

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 (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 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

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.

$\texttt{get_type()} \to \mathsf{str}$

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.

$\texttt{get_type()} \to str$

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.

 $\texttt{get_type()} \to \mathsf{str}$

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.

$get_mass_fraction_at_lower_interface(region: str, component: str) \rightarrow [List[float], List[float]]$

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)

get_mass_fraction_at_upper_interface(region: str, component: str) → [List[float], 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)

 $\texttt{get_mass_fraction_of_component_at_time}(\textit{component: str, time: Union[SimulationTime, float]}) \rightarrow [\texttt{List[float]},\texttt{List[float]}]$

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)

$get_mass_fraction_of_phase_at_time(phase: str, time: Union[SimulationTime, float]) \rightarrow [List[float]]$ 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) → [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*) \rightarrow [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)

get_mole_fraction_of_component_at_time(component: str, time: Union[SimulationTime, float]) → [List[float], 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)

get_mole_fraction_of_phase_at_time(phase: str, time: Union[SimulationTime, float]) \rightarrow [List[float], 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)

$get_position_of_lower_boundary_of_region(region: str) \rightarrow [List[float], List[float]]$

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])

$get_regions() \rightarrow List[str]$

Returns the regions of the diffusion simulation.

Note: Automatically generated regions $(R_{\#})$ are included in the list.

Returns

The region names

$\texttt{get_time_steps()} \rightarrow List[float]$

Returns the timesteps of the diffusion simulation.

Returns

The timesteps [s]

get_total_mass_fraction_of_component(component: str) → [List[float], 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)

 $\texttt{get_total_mole_fraction_of_component}(\textit{component: str}) \rightarrow [\texttt{List[float]},\texttt{List[float]}]$

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)

 $\texttt{get_total_mole_fraction_of_component_in_phase(\mathit{component: str, phase: str}) \rightarrow [\texttt{List[float]}, \texttt{float}], \texttt{float}],$

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

 $\ensuremath{\textbf{phase}}\xspace$ – 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)

 $get_values_of(x_axis: Union[DiffusionQuantity, str], y_axis: Union[DiffusionQuantity, str], plot_condition: Union[PlotCondition, str] = ", independent_variable: Union[IndependentVariable, str] = ") <math>\rightarrow$ [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])

```
get_velocity_of_upper_boundary_of_region(region: str) → [List[float], List[float]]
```

Returns the velocity 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], 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

region – The 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

$\texttt{get_type()} \to \mathsf{str}$

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

$\texttt{get_type()} \to str$

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.

$\texttt{get_type()} \to \mathsf{str}$

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]* =

[])

(1)

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.

get_geometrical_factor() \rightarrow float

Returns the geometrical factor.

Returns

The geometrical factor

$\texttt{get_no_of_points()} \rightarrow \text{int}$

Returns the number of grid points.

Returns

The number of grid points

$\texttt{get_type()} \to str$

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.

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 General Upper Hashin Shtrikman 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

$\texttt{get_type()} \to str$

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

 $\texttt{get_type()} \to str$

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.

 $\texttt{get_type()} \to \mathsf{str}$

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

 $\texttt{get_type()} \to \mathsf{str}$

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.

$get_type() \rightarrow str$

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

 $\texttt{set_max_timestep_allowed_as_percent_of_simulation_time}(max_timestep_allowed_as_percent_of_simulation_time: float = 10.0)$

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

5.1.7 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().

add_poly_command(poly_command: str)

This function has no effect and will be removed in 2024b. If you use it, please remove from your code.

Parameters poly_command -

Returns

This PropertyModelCalculation object

 $calculate(timeout_in_minutes: float = 0.0) \rightarrow PropertyModelResult$

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.

$\texttt{get_argument_default}(\textit{argument_id: str}) \rightarrow \texttt{object}$

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

The detailed description

$\texttt{get_arguments()} \rightarrow Set[str]$

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

$\texttt{get_dynamic_arguments()} \rightarrow \texttt{Set[str]}$

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]

$\texttt{get_model_parameters()} \rightarrow Set[str]$

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_all_poly_commands()

This function has no effect and will be removed in 2024b. If you use it, please remove from your code.

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.

$\texttt{get_result_quantities()} \rightarrow \texttt{Set[str]}$

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(*result_quantity_id: str*) → 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

5.1.8 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]

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

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

$\texttt{get_components()} \rightarrow List[str]$

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

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

 $\ensuremath{\textbf{unit}}$ – 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

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 '='.

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*')

The phase diagram data

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'), *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, ...).

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

$\texttt{get_gibbs_energy_addition_for}(\textit{phase: str}) \rightarrow \texttt{float}$

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

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

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($x_quantity: Union[ThermodynamicQuantity, str], y_quantity:$ $Union[ThermodynamicQuantity, str], sort_and_merge: bool =$ $<math>True) \rightarrow Dict[str, ResultValueGroup]$

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',tity.user_defined_function('HM.T'))

ThermodynamicQuan-

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 '='.

Exampleget_values_grouped_by_quantity_of('T',ThermodynamicQuan-tity.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*')

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 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 *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

$\texttt{get_components()} \rightarrow List[str]$

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.

$\texttt{get_system_data()} \rightarrow \textit{SystemData}$

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

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

 $\textbf{condition} - The \ condition$

Returns

This MaterialToMaterialSingleEquilibriumCalculation object

with_options(options: SingleEquilibriumOptions)

Sets the simulation options.

Parameters

options – The simulation options

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

 $\textbf{condition}-The \ condition$

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().

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.

5.1.9 Package "process_metallurgy"

5.1.9.1 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.

$\texttt{get_composition()} \rightarrow \textit{Dict[str, float]}$

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

$\texttt{get_dependent_component()} \rightarrow str$

Returns the dependent component.

Returns

The dependent component or an empty string if no dependent component is defined

$\texttt{get_elements()} \rightarrow Set[str]$

Returns all elements of the addition.

Returns

The elements

$\texttt{get_id}() \to \text{str}$

Returns the unique ID of the addition.

Returns

The unique ID of the addition

$\texttt{get_temperature()} \to \texttt{float}$

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.

TCOX8 = 2

The database TCOX8.

TCOX9 = 3

The database TCOX9.

$get_name() \rightarrow str$

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

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

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.

5.1.9.2 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]

$get_composition_unit() \rightarrow CompositionUnit$

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 = GasAmountUnit.NORM_CUBIC_METER, composition_unit: GasCompositionUnit = 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

 $\texttt{get_composition_unit()} \rightarrow \textit{GasCompositionUnit}$

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.

get_activity_of_slag(*component: str*, *reference:* ActivityReference = ActivityReference.LIQUID) \rightarrow 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]

get_amount_of_phase_groups() → Dict[PhaseGroup, float]

Returns the amount of each phase group (e.g., for example all liquid slag).

Returns

The amount of the phase groups [kg]

$\texttt{get_amount_of_phases()} \rightarrow \textit{Dict[str, float]}$

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

 $\texttt{get_composition}(composition_unit: CompositionUnit = CompositionUnit.MASS_PERCENT) \rightarrow \texttt{Dict[str, float]}$

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**.

The composition

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

```
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) \rightarrow 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 $get_slag_property()$.

- **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**

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

$\texttt{get_gas_components()} \rightarrow Set[str]$

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

$\texttt{get_oxygen_partial_pressure()} \rightarrow \texttt{float}$

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]

$\texttt{get_stable_phases()} \rightarrow \texttt{Set[str]}$

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

$\texttt{get_temperature()} \rightarrow \texttt{float}$

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

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

5.1.9.3 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.

class tc_python.process_metallurgy.process.**BulkZone**(*density: float, phase_group_to_transfer:* PhaseGroup, name: str)

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]

$get_elements() \rightarrow Set[str]$

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

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

class tc_python.process_metallurgy.process.ContinuousAddition(composition: Dict[str, float], rate:

float, temperature: float = 293.15, composition_unit: CompositionUnit = CompositionUnit.MASS_PERCENT, do_scale: bool = False)

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.

- 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

get_composition_unit() → CompositionUnit

Returns the composition unit used in this addition.

Returns

The composition unit

 $\texttt{get_rate()} \to \texttt{float}$

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.

$get_composition_unit() \rightarrow GasCompositionUnit$

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()*.

$\texttt{get_amount()} \rightarrow List[float]$

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]

$\texttt{get_amount_of_components()} \rightarrow \texttt{Dict[str, List[float]]}$

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

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

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

$\texttt{get_id}() \to str$

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

$\texttt{is_degassing_enabled()} \rightarrow \texttt{bool}$

Returns if degassing is enabled in the zone.

Returns

If degassing is enabled

class tc_python.process_metallurgy.process.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]

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.

- 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**

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]

get_amount_of_elements() → 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]

$\texttt{get_components()} \rightarrow Set[str]$

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

get_composition_of_phase_group(zone: Union[Zone, str], phase_group: PhaseGroup,

composition_unit: CompositionUnit = *CompositionUnit.MASS_PERCENT, composition_type:* CompositionType = CompositionType.COMPONENT) → 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

$\texttt{get_elements()} \rightarrow \texttt{Set[str]}$

Returns all elements present in the simulation.

Returns

The elements

$get_enthalpy() \rightarrow List[float]$

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

```
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) → 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

 $\texttt{get_fraction_of_phases}(\textit{zone: Union[Zone, str], unit: PhaseUnit = PhaseUnit.MASS_FRACTION}) \rightarrow \\ \texttt{Dict[str, List[float]]}$

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

$get_gas_components() \rightarrow Set[str]$

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

get_stable_phases_in_phase_group(zone: Union[Zone, str], phase_group: PhaseGroup) \rightarrow 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*]) → 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]

$\texttt{get_time_points()} \rightarrow List[float]$

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

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]

Returns the kinematic viscosity of a phase in a zone at each time point.

- zone The zone object or the zone name
- **phase** The phase name

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]

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

$get_elements() \rightarrow Set[str]$

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]

$\texttt{get_composition_unit()} \rightarrow \textit{CompositionUnit}$

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.

- 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

$\texttt{get_amount()} \to \texttt{float}$

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.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.

- addition A ContinuousAddition or ContinuousGasAddition
- from_time The start time point [s]
- to_time The end time point [s]

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

get_density() \rightarrow float

Returns the density of the zone

Returns

The density [kg/m**3]

$get_elements() \rightarrow Set[str]$

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

$\texttt{get_phase_group_to_transfer()} \rightarrow \textit{PhaseGroup}$

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

class tc_python.process_metallurgy.process.TransferOfPhaseGroup(phase_group_to_transfer:

PhaseGroup, *source_zone*: Zone)

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

$\texttt{get_transfer_source_zone_id()} \rightarrow str$

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.

- addition A ContinuousAddition or ContinuousGasAddition
- **from_time** The start time point [s]
- **to_time** The end time point [s]

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

$\texttt{get_elements()} \rightarrow \texttt{Set[str]}$

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

is_degassing_enabled() \rightarrow bool

Returns if degassing is enabled in the zone.

Returns

If degassing is enabled

5.1.10 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

$\texttt{get_configuration_as_string()} \to \mathsf{str}$

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.

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()*.

- 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)

Tuple containing the pyvista.Plotter and the pyvista.DataSet mesh object

$\texttt{get_result_file_path()} \rightarrow \textit{str}$

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

class tc_python.am.Automatic

Bases: FileSavingStrategy

An automatic saving strategy.

$get_type() \rightarrow str$

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

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).

$\texttt{get_type()} \to \mathsf{str}$

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 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

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: HeatSource

A conical heat source.

$\texttt{get_type()} \to \textit{str}$

Returns the type of heat source.

Returns

The type

set_absorptivity(absorptivity: float = 60.0)

Sets the absorptivity.

Default: 60%

Parameters absorptivity – The absorptivity [%]

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.**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 = {'focal_point': [0.0, 0.0, 0.0], 'position': [-2, -2, 1],
'viewup': [0, 0, 1]}
```

The default pyvista camera view.

class tc_python.am.DoubleEllipsoidalHeatSource

Bases: HeatSource

A double ellipsoidal heat source.

 $\texttt{get_type()} \rightarrow \textit{str}$

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]

This DoubleEllipsoidalHeatSource object

class tc_python.am.EveryNthTimeStep

Bases: FileSavingStrategy

Saving at every n-th time step.

$\texttt{get_type()} \to \mathsf{str}$

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.

$get_type() \rightarrow str$

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

Bases: HeatSource

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.

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 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, 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 double_ellipsoidal()

A double ellipsoidal heat source.

The default is a Gaussian heat source.

Returns

A new DoubleEllipsoidalHeatSource object

classmethod gaussian()

A Gaussian heat source.

This is the default.

Returns

A new GaussianHeatSource object

$\texttt{classmethod get_calibrated_heatsource_names()} \rightarrow List[str]$

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

Returns the file path of the calibrated heat source.

Parameters

optimized_heatsource_name – The name of the calibrated heat source

The file path

class tc_python.am.KeyholeModel

Bases: object

A model for an "analytic" keyhole in the AM calculation.

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)

Creates material properties from the result of a Scheil calculation.

Parameters

result - The Scheil result to create material properties from

Returns

A new ScheilMaterialProperties object

$\texttt{classmethod get_all_library_names()} \rightarrow List[str]$

Returns a list with the names of all material libraries available on disk.

Returns

A list with the names of all material libraries

$get_average_material_property_in_range(material_property_enum: MaterialProperty, from_zone: Zone, to_zone: Zone) \rightarrow float$

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

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]

$\texttt{get_liquidus_temperature()} \rightarrow \texttt{float}$

Returns the liquidus temperature for the material.

Returns

The liquidus temperature [K]

$\texttt{get_name()} \to \mathsf{str}$

Returns the name of the library.

Returns

The name of the library

get_smoothed_values_for_material_property(*material_property_enum*: 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

$\texttt{get_solidification_temperature()} \rightarrow \texttt{float}$

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]

ENTHALPY_PER_KG = 10

Enthalpy [J/kg]

$ENTHALPY_PER_MOLE = 2$

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

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

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

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

class tc_python.am.ProbeCoordinate(x: float, y: float, z: float)

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.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

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)

Bases: MaterialProperties

Material properties created from the result of a Scheil calculation.

class tc_python.am.SingleTrackScanningStrategy

Bases: ScanningStrategy

A single track scanning strategy.

$get_type() \rightarrow str$

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.

get_heat_affected_zone_depth() → float

Returns the depth of the heat affected zone.

Returns

The depth of the heat affected zone [m]

get_heat_affected_zone_length() \rightarrow float

Returns the length of the heat affected zone.

Returns

The heat affected zone length [m]

get_heat_affected_zone_width() → 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.

The width of the keyhole [m]

$\texttt{get_meltpool_depth()} \to \texttt{float}$

Returns the meltpool depth.

Returns

The meltpool depth [m]

$\texttt{get_meltpool_length()} \rightarrow \texttt{float}$

Returns the meltpool length.

Returns

The meltpool length [m]

get_meltpool_width() → 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.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

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])

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.

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_properties – 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).

$\texttt{get_type()} \to str$

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.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

5.2 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

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

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_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

$\texttt{get_elements_in_system()} \rightarrow List[str]$

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

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

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

get_species_object(species_name: str) \rightarrow Species

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

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 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

5.3 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() → 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]

get_molar_mass() \rightarrow float

Returns the molar mass of the element.

Returns

The molar mass [g/mol]

$\texttt{get_name()} \to \textit{str}$

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

$\texttt{get_sublattices()} \rightarrow \texttt{List}[\textit{Sublattice}]$

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

$\texttt{has_diffusion_data()} \rightarrow \texttt{bool}$

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

$is_gas() \rightarrow bool$

Returns if the phase is a gas phase.

Returns

If the phase is a gas phase

 $\texttt{is_ionic_liquid()} \rightarrow \texttt{bool}$

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

$\textbf{is_solid()} \rightarrow bool$

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

5.4 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) \rightarrow

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

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

get_database_info(*database_short_name: str*) → str

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

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

 $select_thermodynamic_and_kinetic_databases_with_elements(thermodynamic_db_name: str, kinetic_db_name: str, list_of_elements: List[str]) \rightarrow MultiDatabaseSystemBuilder$

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*.

5.5 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 [m²/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.

<code>classmethod time()</code> \rightarrow Time

Creates a quantity representing the time [s].

classmethod total_mass_fraction_of_component(component: str) \rightarrow

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.

$\texttt{classmethod velocity_of_upper_boundary_of_region(\mathit{region: str})} \rightarrow$

VelocityOfUpperBoundaryOfRegion

Creates a quantity representing the velocity of upper boundary of a region [m/s].

Parameters

 $\ensuremath{\textbf{region}}\xspace - The name of the region$

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.

$\texttt{classmethod heat_per_gram()} \rightarrow HeatPerGram$

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.

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.

$\texttt{classmethod latent_heat_per_gram()} \rightarrow LatentHeatPerGram$

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.

$\texttt{classmethod mass_fraction_of_a_solid_phase(\textit{phase: str}) \rightarrow \texttt{MassFractionOfASolidPhase}}$

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. classmethod mass_fraction_of_all_solid_phases() \rightarrow MassFractionOfAllSolidPhase Creates a quantity representing the total mass fraction of all solid phases. Returns A new MassFractionOfAllSolidPhase 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. $\texttt{classmethod molar_volume_of_system}() \rightarrow MolarVolumeOfSystem}$ Creates a quantity representing the molar volume of the system [m³/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. $classmethod mole_fraction_of_all_liquid() \rightarrow MoleFractionOfAllLiquid$ Creates a quantity representing the total molar fraction of all the liquid phase. Returns A new MoleFractionOfAllLiquid object. classmethod mole_fraction_of_all_solid_phases() \rightarrow 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.

$\texttt{classmethod temperature()} \rightarrow Temperature$

Creates a quantity representing the temperature [K].

Returns

A new Temperature 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.

 $\texttt{classmethod volume_fraction_of_all_liquid()} \rightarrow VolumeFractionOfAllLiquid$

Creates a quantity representing the total volume fraction of all the liquid phase.

Returns

A new VolumeFractionOfAllLiquid object.

$classmethod \ volume_fraction_of_all_solid_phases() \rightarrow VolumeFractionOfAllSolidPhases() \rightarrow VolumeFractionOfAllPhases() \rightarrow VolumeFractionOfAllPhases() \rightarrow Volum$

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 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 composition_of_phase_as_mole_fraction(phase: str, component: str = 'All') \rightarrow CompositionOfPhaseAsMoleFraction
```

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.

 $\texttt{classmethod composition_of_phase_as_weight_fraction}(\textit{phase: str, component: str}) \rightarrow \texttt{}$

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 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 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 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 system_size() \rightarrow SystemSize

Creates a quantity representing the system size [mol].

Returns

A new SystemSize object.

$\texttt{classmethod temperature()} \rightarrow Temperature$

Creates a quantity representing the temperature [K].

Returns

A new Temperature object.

classmethod tracer_diffusion_coefficient(phase: str, diffusing_element: str) \rightarrow TracerDiffusionCoefficient

Creates a quantity representing tracer diffusion coefficient of a phase [m²/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.

5.6 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

 $\texttt{get_x()} \rightarrow \texttt{List[float]}$

Accessor for the x-values :return the x values

 $\texttt{get}_\texttt{y}() \to \texttt{List}[\texttt{float}]$

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

5.7 Module "propertymodel_sdk"

```
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

$\texttt{get_argument_ids()} \rightarrow \texttt{Set[str]}$

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, \dots)

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() \rightarrow 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]

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]

$\texttt{get_mole_fractions()} \rightarrow Dict[str, float]$

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]

$\texttt{get_mole_percents()} \rightarrow Dict[str, float]$

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]

$\texttt{get_temperature()} \to \texttt{float}$

Obtains the current temperature from the UI.

Returns

The temperature [K]

$\texttt{get_ui_boolean_value}(\textit{component_id: str}) \rightarrow \texttt{bool}$

Obtains the value from the specified checkbox UI component.

Parameters component_id - Id of the checkbox

Returns

The setting of the checkbox

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

component_id – Id of the list UI component

Returns

The selected entry

$\texttt{get_ui_string_value}(\textit{component_id: str}) \rightarrow \texttt{str}$

Obtains the selected entry from a UI component text field.

Parameters

 $\textbf{component_id} - Id \ of \ the \ string \ UI \ component$

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)

 $\label{eq:get_description} \ensuremath{\mathsf{get_description}}\xspace) \to \operatorname{str}$ Obtains the description of the quantity.

Returns Description of the quantity

$\texttt{get_id}() \to \text{str}$

Obtains the id of the quantity.

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

STRENGTH_QUANTITY = 8

A strength quantity

SURFACE_ENERGY_QUANTITY = 3

A surface energy quantity

TEMPERATURE_QUANTITY = 1

A temperature quantity

$TIME_QUANTITY = 4$

A time 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

$get_setting() \rightarrow bool$

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

get_description() \rightarrow str

Obtains the additional description of the component.

Returns

Additional description of the component

$\texttt{get_id}() \to \text{str}$

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

class tc_python.propertymodel_sdk.UIConditionListComponent(component_id: str, name: str,

description: str)

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

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)

$\texttt{get_selected_entry()} \rightarrow str$

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

any_marker_setting: bool = False)

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

$\texttt{get_default_phase()} \rightarrow \textit{str}$

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: 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

$\texttt{get_value()} \rightarrow \textit{str}$

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

class tc_python.propertymodel_sdk.UIStringComponent(component_id: str, name: str, description: str,

string: 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

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

$\texttt{get_value()} \to \mathsf{str}$

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

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

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) $\rightarrow ResultQuantity$

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

tc_python.propertymodel_sdk.create_phase_list_ui_component(component_id: str, name: str,

description: str, default_phase: str = ", any_marker: bool = False) \rightarrow UIPhaseListComponent 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_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

 $\texttt{tc_python.propertymodel_sdk.create_string_ui_component(component_id: str, name: str, description: str, string: str}) \rightarrow UIStringComponent$

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

 $\texttt{tc_python.propertymodel_sdk.create_surface_energy_quantity(\textit{quantity_id: str, description: str}) \rightarrow \textit{ResultQuantity} \rightarrow \textit{ResultQuantity}$

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) $\rightarrow ResultQuantity$

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

 $\texttt{tc_python.propertymodel_sdk.create_time_quantity}(\textit{quantity_id: str, description: str}) \rightarrow \textit{ResultQuantity}(\textit{quantity_id: str, description: str, description: str_{description: str_{description: str_{description:$

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

5.8 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)

5.9 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

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

$\texttt{get_system_function_names()} \rightarrow List[str]$

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() → List[TemperatureInterval]

Returns the list of all defined intervals.

Returns

The defined temperature intervals

$\texttt{get_lower_temperature_limit()} \rightarrow \texttt{float}$

Returns the lower temperature limit.

Returns

The lower temperature limit in K

$get_name() \rightarrow str$

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(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

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

get_expression() \rightarrow str

Returns the function expression of this temperature interval.

Returns

The temperature function expression

get_upper_temperature_limit() \rightarrow 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

CHAPTER

TROUBLESHOOTING

This section provides an FAQ for common problems that occur when using TC-Python.

6.1 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 = '2024a'
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)
```

(continued from previous page)

```
'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 + ')')

print('')
print('Url of license server: (if license server is NO-NET, you need a local license
→file)')
try:
   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)')
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__)
with tc_python.TCPython() as session:
   print('')
   print('Lists the databases: (should be a complete list of the installed databases.
→that you have license for or do not require license)')
   print(session.get_databases())
```

6.2 "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*.

6.3 "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* (has transient dependency *six*)
- *pyvista* (*optional*, *only for Additive Manufacturing result plotting and data extraction*, it has a number of transient dependencies)
- 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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