Asim Documentation

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Matěj Laitl

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Asim Project

1.1 Launching

```
Simple simulation can be starting by calling:
```

```
python -c 'import simulation.simple as s; s.main()
```

Assimilation can be started using

```
python -c 'import assimilation.twin as t; t.main()
```

Calling something like *python simulation/simple.py* doesn't work for Cython build because it circumvents binary version of the simulation.simple module. The main() methofs may take optional parameters, study the sources.

1.2 Cythoning

For dramatic performance improvements you can Cythonize Python modules into C and then compile them forming binary Python modules. You need to have recent Cython (0.17 beta3 or newer) and a compiler. Cythoning is performed by calling python setup.py build_ext -i. Every time you change source files you need to re-Cython. Beware, Python prefers binary modules to .py modules, because of that command python setup.py uncython to clean any binary modules and generated C.

Dispersion Model

Module with interfaces (empty classes) for core dispersion model objects.

class dispmodel.iface.DispersionModel

Bases: object

Model of propagation of radionuclides in atmosphere.

concentration_at()

Compute radioactive concentration at given location (Location (page 3)) and model's current time. TODO: units.

Returns view on 1D array with per-nuclide concentration. You may *not* modify values in the view. The view may become invalid when this or other method is called on this instance again

deposition_at()

Compute combined dry and wet radioactive deposition at given location (Location (page 3)) and model's current time. TODO: units.

Returns view on 1D array with per-nuclide deposition. You may *not* modify values in the view. The view may become invalid when this or other method is called on this instance again

dose_at()

Compute radioactive dose at given location (Location (page 3)) and model's current time in Sieverts.

Returns view on 1D array with per-nuclide dose. You may *not* modify values in the view. The view may become invalid when this or other method is called on this instance again

propagate()

Propagate model to next time step.

Parameters

- meteo_model (MeteoModel (page 4)) model of weather the calculation should use
- source_model (SourceModel (page 4)) model of the source of the release the calculation should use

class dispmodel.iface.Location

Bases: object

Describes a point in 3D space.

class dispmodel.iface.MeteoModel

Bases: object

Model of weather.

dispersion_xy()

Return horizonal dispersion coefficient for a puff that is currently at location *loc*, time *time* and has flewn *total* distance meters in total.

dispersion_z()

Return vertical dispersion coefficient for a puff that is currently at location *loc*, time *time* and has flewn *total distance* meters in total.

mixing_layer_height_at()

Return height of the mixing layer in meters at given location and time.

Parameters

- loc (Location (page 3)) where the height is to be computed
- time (int) time in seconds since start of the simulation

Return type float

wind_direction_at()

Return direction of the wind in radians at given location and time.

Parameters

- loc (Location (page 3)) where the direction is to be computed
- time (int) time in seconds since start of the simulation

Return type float

wind_speed_at()

Return speed of the wind in m/s at given location and time.

Parameters

- loc (Location (page 3)) where the speed is to be computed
- time (int) time in seconds since start of the simulation

Return type float

class dispmodel.iface.Nuclide

Bases: object

Nuclide describes properties of one radionuclide.

class dispmodel.iface.SourceModel

Bases: object

Model of the source of an atmospheric radioactive release.

inventory()

Return an array of radionuclides this release contains.

Return type array of Nuclides (page 4)

location()

Return location of the release. All three coordinates are to be used.

Return type Location (page 3)

release_rate()

Return release rate in Bq/s of all radionuclides at given time. Rate of radionuclide inventory()[i] (page 4) is release_rate(time)[i] (page 5).

Return type array of floats

 ${\bf class}$ dispmodel.iface._memoryviewslice

Bases: dispmodel.iface.memoryview

Nuclide Database

A simple database of radionuclides.

To be used as:

```
>>> import nuclides
>>> argon = nuclides.db["Ar-41"]
>>> argon.half_life
6560.0
```

class dispmodel.nuclides._memoryviewslice Bases: dispmodel.nuclides.memoryview

Puff Model (Dispersion Model Implementation)

Implementation of the dispersion model based on the concept of puffs.

class dispmodel.puffmodel.Puff

Bases: dispmodel.iface.DispersionModel (page 3)

One puff. Makes sense only as a part of PuffModel (page 9).

Parameters

- time_step (int) number of seconds the integration step lasts. propagate() (page 3) advances time_step seconds
- inventory (array of Nuclide (page 4) objects) radionuclides present in this release
- Q (array of floats) per-nuclide initial activities TODO: in what units?
- loc (Location (page 3)) initial location of the puff; Puff references it (no copy)
- time (int) total time in seconds since start of the simulation

dose_at()

Computes dose at location loc using m/mju method

${\bf class} \ {\tt dispmodel.puffmodel.PuffModel}$

```
Bases: dispmodel.iface.DispersionModel (page 3)
```

Implementation of the dispersion model based on the concept of puffs (page 9).

Parameters

- time step (int) propagate() (page 3) advances time_step seconds
- puff_sampling_step (int) inteval between 2 puff releases; must be multiple of time_step. Source model's (page 4) release_rate() (page 5) is queried in puff sampling step-long intervals
- release loc (Location (page 3)) location where the puffs get released
- source model (SourceModel (page 4)) source term of the release

class dispmodel.puffmodel._memoryviewslice

 $Bases: \verb"dispmodel.puffmodel.memoryview"$

Simulation of the Puff Model

Run a simulation of an atmospheric radioactive realease using the puff model

```
class simulation.simple.PasquillsMeteoModel
```

Bases: dispmodel.iface.MeteoModel (page 4)

Abstract meteo model based on Pasquills stability category

Parameters stability_category (int) - Pasquills statiblity category A=0 ... F=5

 ${\bf class} \ {\tt simulation.simple.SimulatedSourceModel}$

Bases: dispmodel.iface.SourceModel (page 4)

Source model that releases static pre-defined ammount of radionuclides.

Parameters

- time step (int) inverval between rate changes
- rates (*list*) total activities per while time slots

${\bf class}$ simulation.simple.StaticMeteoModel

Bases: simulation.simple.PasquillsMeteoModel (page 11)

Weather model that returns pre-computed numbers stored in .mat file.

Parameters

- matfile (*string*) path to Matlab .mat file with pre-computed grid of wind speed and direction
- stability category (int) Pasquills statiblity category A=0 ... F=5

```
_bilinear_interp()
```

Bilinear interpolation.

 ${\bf class} \ {\tt simulation.simple._memoryviewslice}$

Bases: simulation.simple.memoryview

Internal class for passing memoryview slices to Python

simulation.simple.read_receptor_locations()

Loads list of receptors from a file

Returns list of Locations (page 3) with receptor locations

Asimilation Twin Experiment

```
Atmospheric radioactive release twin experiment using simulation.simple (page 11) and
dispmodel.puffmodel (page 9)
class assimilation.twin.AssimilationMeteoModel
     Bases: simulation.simple.PasquillsMeteoModel (page 11)
     Meteorologic model used for assimilation of wind speed and direction.
     Construct assimilation model.
          Parameters stability category (int) – Pasquills statiblity category A=0 ...
class assimilation.twin.ObservationCPdf
     Bases: pybayes.pdfs.CPdf
     (wind speed, wind direction) is not a complete state - it needs to include puff locations,
     total flewn distances etc. Therefore ObservationCPdf references PuffModelEmpPdf to
     acces full state through its models attribute.
     eval_log()
          Evalate cpdf in point x given condition cond.
              Parameters
                  • x – observation vector (wind speed, wind direction, doses measured at
                   receptors)
                  • cond – state vector (at, bt)
class assimilation.twin.TransitionCPdf
     Bases: pybayes.pdfs.CPdf
     TODO: This can be now made ProdCPdf
     sample()
          Sample from this CPdf.
              Parameters cond – condition: previous a t, b t
class assimilation.twin._memoryviewslice
     Bases: assimilation.twin.memoryview
```

Math Wrappers (Cythonization helpers)

```
class support.mathwrapper._memoryviewslice
    Bases: support.mathwrapper.memoryview
    Internal class for passing memoryview slices to Python
support.mathwrapper.iadd_vv()
    Hack for Cython - in-place addition of 2 vectors, same as lhs += rhs
support.mathwrapper.imult_vs()
    Hack for Cython - in-place element-wise multiplication of 2 vectors, same as lhs *= rhs
support.mathwrapper.imult_vv()
    Hack for Cython - in-place element-wise multiplication of 2 vectors, same as lhs *= rhs
support.mathwrapper.matrix()
    Hack for Cython - creates a suitable container for a matrix of C doubles (Python floats)
support.mathwrapper.sum_v()
    Hack for Cython - sums all elements of vector summand
support.mathwrapper.vector()
    Hack for Cython - creates a suitable container for a vector of C doubles (Python floats)
```

Graph Plotters

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