
emodpy-generic

Institute for Disease Modeling

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emodpy-generic is a collection of Python scripts and utilities created to streamline user interactions with EMOD and idmtools for modeling generic diseases. Much of the functionality is inherited from the [emod_api](#) package and [emodpy](#) package.

Additional information about how to use idmtools can be found at in [Welcome to idmtools](#). Additional information about EMOD HIV parameters can be found in [EMOD parameter reference](#).

If you have questions, see [Frequently asked questions](#) for general questions about all emodpy functionality or *Frequently asked questions* for functionality specific to emodpy-generic.

See [Welcome to idmtools](#) for a diagram showing how idmtools and each of the related packages are used in an end-to-end workflow using EMOD as the disease transmission model.

EMODPY-GENERIC INSTALLATION

Follow the steps below to install emodpy-generic.

1.1 Prerequisites

First, ensure the following prerequisites are met.

- Windows 10 Pro or Enterprise, Linux, or Mac
- Python 3.9 64-bit (<https://www.python.org/downloads/release>)
- A file that indicates the pip index-url:
 - Windows
 - Linux

In C:\Users\Username\pip\pip.ini, containing the following:

```
[global]
index-url = https://packages.idmod.org/api/pypi/pypi-production/simple
```

In \$HOME/.config/pip/pip.conf, containing the following:

```
[global]
index-url = https://packages.idmod.org/api/pypi/pypi-production/simple
```

1.2 Installation instructions

1. Open a command prompt and create a virtual environment in any directory you choose. The command below names the environment “v-emodpy-generic”, but you may use any desired name:

```
python -m venv v-emodpy-generic
```

2. Activate the virtual environment:

- Windows
- Linux

Enter the following:

```
v-emodpy-generic\Scripts\activate
```

Enter the following:

```
source v-emodpy-generic/bin/activate
```

3. Install emodpy-generic packages:

```
pip install emodpy-generic
```

If you are on Linux, also run:

```
pip install keyrings.alt
```

4. When you are finished, deactivate the virtual environment by entering the following at a command prompt:

```
deactivate
```


API REFERENCE

2.1 emodpy_generic package

The emodpy-generic module is intended to house scripts and tools that enable disease modelers to work more easily with the IDM EMOD Generic model.

2.1.1 Subpackages

emodpy_generic.demographics package

Submodules

emodpy_generic.demographics.GenericDemographics module

```
class emodpy_generic.demographics.GenericDemographics(nodes,  
                                                    idref='Gridded  
world  
grump2.5arcmin',  
                                                    base_file=None)
```

Bases: `emod_api.demographics.Demographics.Demographics`

This class is derived from emod_api.demographics' Demographics class so that we can set certain defaults for Generic in construction. As we add other disease types, the generalizations and specifics will become clearer.

```
emodpy_generic.demographics.GenericDemographics.fromBasicNode(lat=0, lon=0,  
                                                             pop=1000000.0,  
                                                             name=1,  
                                                             forced_id=1)
```

This function creates a single-node GenericDemographics instance from the params you give it.

Parameters

- **lat** – latitude (not really used)
- **lon** – longitude (not really used)
- **pop** – population. Defaults to 1 million.
- **name** – node name (not really used)
- **forced_id** – node id (not really used)

Returns GENERIC_SIM demographics instance which can be customized and/or written to file.

emodpy_generic.interventions package

The below imports let you do stuff like: `from emodpy_generic.interventions import * a = SimpleSIA()`

If you want to access all the iv function creates directly without scoping/qualifying them.

Submodules

emodpy_generic.interventions.complex_import module

```
emodpy_generic.interventions.complex_import.ComplexImportationEvent (camp,  
                                                                    dips=[0.00012919896640826872,  
                                                                    durs=[100000],  
                                                                    timestep=1,  
                                                                    nods=[])
```

Seed infection (over time) in the modeled population.

Be careful when configuring import pressure in multi-node simulations. `Daily_Import_Pressures` defines a rate of per-day importation for each node that the intervention is distributed to. In a 10 node simulation with `Daily_Import_Pressures = [0.1, 5.0]`, the total importation rate summed over all nodes will be 1/day and 50/day during the two time periods. You must divide the per-day importation rates by the number of nodes. Note that there is no control over the ages of the individuals imported, or their monte-carlo weight, like you might find with the Outbreak intervention.

Parameters

- **dips** – `DailyImportPressures`. An array of rates of per-day importation (per-node).
- **durs** – An array of durations over which to apply import pressure. Goes with `dips`.
- **timestep** – WHEN: Day to start campaign event.
- **nods** – WHERE: List of nodes at which infections will be seeded.

Returns Campaign event that can be added to campaign.

emodpy_generic.interventions.emergence module

```
emodpy_generic.interventions.emergence.basicEmergenceEvent (camp,    genome=0,  
                                                            clade=0, start_day=1,  
                                                            nods=[])
```

Seed infection at a particular time by importing a new infected individual.

Note that this function imports 20 individuals age 5. This could be made configurable but is not yet because this was created with a particular use case in mind. All other values are set to schema defaults.

Parameters

- **genome** – Genome of new infection.
- **clade** – Clade of new infection.
- **start_day** – WHEN: Day to start campaign event.
- **nods** – WHERE: List of nodes at which infections will be seeded.

Returns Campaign event that can be added to campaign.

`emodpy_generic.interventions.emergence.basicSIAEvent` (*camp*, *cov=1.0*, *genome=0*,
clade=0, *start_day=1*,
nods=[])

Seed infection at a particular time by infected existing individuals (or agents really) in the simulation.

Note that the targeted ages are 0-5 years, not configurable without code change.

Parameters

- **clade** – Clade of new infection.
- **genome** – Genome of new infection.
- **cov** – WHO. Percentage of population to infect (given other targeting constraints).
- **start_day** – WHEN: Day to start campaign event.
- **nods** – WHERE: List of nodes at which infections will be seeded.

Returns Campaign event that can be added to campaign.

`emodpy_generic.interventions.emergence.new_intervention_as_file` (*timestep*, *file-*
name=None)

This function mostly exists for testing so one can exercise the functionality in a single function call.

emodpy_generic.interventions.polio_vaccine module

`emodpy_generic.interventions.polio_vaccine.SimpleIPV1` (*camp*, *timestep=1*, *cover-*
age=1.0)

IPV (round 1) Campaign Event

AgeAtVaccInDays is hard-coded to 270.

Parameters

- **coverage** – WHO?
- **timestep** – WHEN?

Returns Campaign event that can be added to campaign.

`emodpy_generic.interventions.polio_vaccine.SimpleIPV2` (*camp*, *timestep=1*, *cover-*
age=1.0)

IPV (round 1) Campaign Event

AgeAtVaccInDays is hard-coded to 270.

Parameters

- **coverage** – WHO?
- **timestep** – WHEN?

Returns Campaign event that can be added to campaign.

`emodpy_generic.interventions.polio_vaccine.new_intervention_as_file` (*timestep*,
file-
name=None)

This function mostly exists for testing so one can exercise the functionality in a single function call.

emodpy_generic.interventions.simple_sia module

emodpy_generic.interventions.simple_sia.**SimpleSIA**(*camp*, *Event_Name='SIA'*,
timestep=1, *Target_Age_Min=0.75*,
Target_Age_Max=5.0, *Cover-*
age=1.0, *nods=[]*)

Create an intervention that represents a highly-effective acquisition-blocking vaccination.

Parameters

- **Event_Name** –
- **timestep** – WHEN to distribute the intervention.
- **Target_Age_Min** – WHO to distribute the intervention to (lower age bound).
- **Target_Age_Max** – WHO to distribute the intervention to (upper age bound).
- **Coverage** – WHO to distribute the intervention to (%age of population).
- **nods** – WHERE to distribute (list of nodes).

Returns Campaign event that can be added to campaign.

emodpy_generic.interventions.simple_sia.**new_intervention_as_file**(*timestep*, *file-*
name=None)

This function mostly exists for testing so one can exercise the functionality in a single function call.

FREQUENTLY ASKED QUESTIONS

As you get started with `emodpy-generic`, you may have questions. The most common questions will be added below. For questions related to functionality in related packages, see the following documentation:

- [Frequently asked questions for EMOD](#)
- [Frequently asked questions for idmtools](#)
- [Frequently asked questions for emod-api](#)
- [Frequently asked questions for emodpy](#)

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