epiworld

0.0-1

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Chapter 1

Main Page

1.1 epiworld

This C++ template-header-only library provides a general framework for epidemiologic simulation. The main features of the library are:

- 1. Four key classes: Model, Person, Tool, and Virus.
- 2. The model features a social networks of Persons.
- 3. Persons can have multiple Tools as a defense system.
- 4. Tools can reduce contagion rate, transmissibility, death rates, and improve recovery rates.
- 5. Viruses can mutate (generating new variants).
- 6. Models can feature multiple states, e.g., HEALTHY, SUSCEPTIBLE, etc.
- 7. Models can have an arbitrary number of parameters.
- 8. **REALLY FAST** About 6.5 Million person/day simulations per second.

1.2 Hello world

Here is a simple SIRS model implemented with

```
#include "../include/epiworld/epiworld.hpp"
using namespace epiworld;
int main()
{
          // Creating a model
          Model<> model;
          // Adding the tool and virus
          Virus<> virus("covid 19");
          virus.set_post_immunity(1.0);
          model.add_virus_n(virus, 5);

          Tool<> tool("vaccine");
          model.add_tool(tool, .5);
          // Generating a random pop
          model.population_smallworld(100000);
          // Initializing setting days and seed
          model.init(100, 123);
```

2 Main Page

```
// Running the model
model.run();
model.print();
```

And you should get something like the following:

Running the model...

```
SIMULATION STUDY
Population size
               : 100000
Days (duration)
               : 100 (of 100)
Number of variants : 1
Last run elapsed t : 280.00ms
Rewiring
               : off
Virus(es):
 - covid 19 (baseline prevalence: 5 seeds)
Tool(s):
  vaccine (baseline prevalence: 50.00%)
Model parameters:
- Total healthy (S) : 99995 -> 97390
- Total recovered (S) : 0 -> 2554
- Total infected (I)
                         5 -> 56
                         0 -> 0
 - Total removed (R)
(S): Susceptible, (I): Infected, (R): Recovered
```

Which took about 0.280 seconds.

1.2.1 **Tools**

1.2.2 Contagion

Susceptible individuals can acquire a virus from any of their infected connections. The probability that susceptible individual i gets the virus v from individual j depends on how three things:

- 1. The transmissibility of the virus, ,
- 2. The contagion reduction factor of i, , and
- 3. The host's transmission reduction factor, .

The last two are computed from and 's tools. Ultimately, the probability of getting virus \$v\$ from equals:

Nonetheless, the default behavior of the simulation model is to assume that individuals can acquire only one disease at a time, if any. This way, the actual probability is:

The latter is calculated using Bayes' rule

Where

This way, viruses with higher transmissibility will be more likely to be acquired when competing with other variants.

Chapter 2

Class Index

2.1 Class List

Here are the classes, structs, unions and interfaces with brief descriptions:

| Action< TSeq > | |
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| Action data for update an agent | 5 |
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| DataBase < TSeq > | |
| Statistical data about the process | 8 |
| LFMCMC< TData > | |
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| Model < TSeq > | |
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| Person < TSeq > | |
| Person (agents) | 19 |
| PersonTools < TSeq > | |
| List of tools available for the individual to | 20 |
| PersonViruses < TSeq > | |
| Set of viruses in host | 21 |
| Progress | |
| A simple progress bar | 22 |
| Queue < TSeq > | |
| Controls which agents are verified at each step | 23 |
| RandGraph | 23 |
| Tool< TSeq > | |
| Tools for defending the host against the virus | 24 |
| UserData < TSeq > | |
| Personalized data by the user | 25 |
| vecHasher< T > | |
| Vector hasher | 27 |
| Virus< TSeq > | |
| Virus | 28 |

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Chapter 3

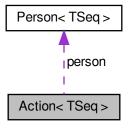
Class Documentation

3.1 Action < TSeq > Struct Template Reference

Action data for update an agent.

#include <config.hpp>

Collaboration diagram for Action < TSeq >:



Public Member Functions

Action (Person< TSeq > *person_, VirusPtr< TSeq > virus_, ToolPtr< TSeq > tool_, epiworld_fast_uint virus_idx_, epiworld_fast_uint new_status_, epiworld_fast_int queue_, Action← Fun< TSeq > call_)

Construct a new Action object.

Public Attributes

- Person< TSeq > * person
- VirusPtr< TSeq > virus
- ToolPtr< TSeq > tool
- epiworld_fast_uint virus_idx
- epiworld_fast_uint tool_idx
- · epiworld_fast_uint new_status
- epiworld_fast_int queue
- ActionFun< TSeq > call

3.1.1 Detailed Description

```
template < typename TSeq > struct Action < TSeq >
```

Action data for update an agent.

Template Parameters

3.1.2 Constructor & Destructor Documentation

3.1.2.1 Action()

Construct a new Action object.

All the parameters are rather optional.

Parameters

| person_ | Person over who the action will happen |
|-----------|--|
| virus_ | Virus to add |
| tool_ | Tool to add |
| virus_idx | Index of virus to be removed (if needed) |
| tool_idx | Index of tool to be removed (if needed) |
| new_← | Next status |
| status_ | |
| queue_ | Efect on the queue |
| call_ | The action call (if needed) |

The documentation for this struct was generated from the following file:

• include/epiworld/config.hpp

3.2 AdjList Class Reference

Public Member Functions

AdjList (const std::vector< unsigned int > &source, const std::vector< unsigned int > &target, bool directed, int min_id=-1, int max_id=-1)

Construct a new Adj List object.

- void read_edgelist (std::string fn, int skip=0, bool directed=true, int min_id=-1, int max_id=-1)
- std::map< unsigned int, unsigned int > operator() (unsigned int i) const
- void print (unsigned int limit=20u) const
- · unsigned int get_id_max () const
- unsigned int get_id_min () const
- size_t vcount () const
- size_t ecount () const
- std::map< unsigned int, std::map< unsigned int, unsigned int > > & get_dat ()
- · bool is directed () const

3.2.1 Constructor & Destructor Documentation

3.2.1.1 AdjList()

Construct a new Adj List object.

It will create an adjacency list object with maxid - minid + 1 nodes. If min_id and max_id are not specified (both < 0), then the program will try to figure them out automatically by looking at the range of the observed ids.

Parameters

| source | Unsigned int vector with the source |
|----------|--------------------------------------|
| target | Unsigned int vector with the target |
| directed | Bool true if the network is directed |
| min_id | int min id. |
| max_id | int max id. |

The documentation for this class was generated from the following files:

- · include/epiworld/adjlist-bones.hpp
- · include/epiworld/adjlist-meat.hpp

3.3 DataBase < TSeq > Class Template Reference

Statistical data about the process.

#include <database-bones.hpp>

Public Member Functions

- DataBase (int freg=1)
- void record_variant (Virus < TSeq > &v)

Registering a new variant.

- void record_tool (Tool < TSeq > &t)
- void set_seq_hasher (std::function< std::vector< int >(TSeq)> fun)
- void set_model (Model < TSeq > &m)
- Model < TSeq > * get_model ()
- void record ()
- const std::vector< TSeq > & get_sequence () const
- const std::vector< int > & get_nexposed () const
- · size t size () const
- void **write_data** (std::string fn_variant_info, std::string fn_variant_hist, std::string fn_tool_info, std::string fn tool_hist, std::string fn_toal_hist, std::string fn_transmission, std::string fn_transmission) const
- void record_transmission (int i, int j, int variant)
- size_t get_nvariants () const
- · void reset ()
- void set_user_data (std::vector< std::string > names)
- void add_user_data (std::vector< epiworld_double > x)
- void **add_user_data** (unsigned int j, epiworld_double x)
- UserData < TSeq > & get_user_data ()

Get recorded information from the model

Parameters

what std::string, The status, e.g., 0, 1, 2, ...

Returns

In get_today_total, the current counts of what.

In get_today_variant, the current counts of what for each variant.

In get_hist_total , the time series of what

In get_hist_variant, the time series of what for each variant.

In get_hist_total_date and get_hist_variant_date the corresponding dates

- int get_today_total (std::string what) const
- int get_today_total (epiworld_fast_uint what) const
- void get today total (std::vector < std::string > *status=nullptr, std::vector < int > *counts=nullptr) const
- void get_today_variant (std::vector < std::string > &status, std::vector < int > &id, std::vector < int > &counts) const
- void get_hist_total (std::vector< int > *date, std::vector< std::string > *status, std::vector< int > *counts) const
- void get_hist_variant (std::vector< int > &date, std::vector< int > &id, std::vector< std::string > &status, std::vector< int > &counts) const

Friends

- class Model < TSeq >
- void default_add_virus (Action< TSeq > &a, Model< TSeq > *m)
- void default_add_tool (Action < TSeq > &a, Model < TSeq > *m)
- void default_rm_virus (Action < TSeq > &a, Model < TSeq > *m)
- void default_rm_tool (Action < TSeq > &a, Model < TSeq > *m)

3.3.1 Detailed Description

```
template<typename TSeq> class DataBase< TSeq>
```

Statistical data about the process.

Template Parameters

3.3.2 Member Function Documentation

3.3.2.1 record_variant()

Registering a new variant.

Parameters

Pointer to the new variant. Since variants are originated in the host, the numbers simply move around.From the parent variant to the new variant. And the total number of infected does not change.

The documentation for this class was generated from the following files:

- · include/epiworld/database-bones.hpp
- include/epiworld/database-meat.hpp

3.4 LFMCMC< TData > Class Template Reference

Likelihood-Free Markov Chain Monte Carlo.

```
#include <1fmcmc.hpp>
```

Public Member Functions

- void run (VEC(epiworld_double) param_init, size_t n_samples_, epiworld_double epsilon_)
- · LFMCMC (TData & observed data)
- void set_observed_data (TData &observed_data_)
- void set_proposal_fun (FUN< void(VEC(epiworld_double)&, LFMCMC< TData > *)> fun)
- void set_simulation_fun (FUN< TData(VEC(epiworld_double)&, LFMCMC< TData > *)> fun)
- void set_summary_fun (FUN< VEC(epiworld_double)(TData &, LFMCMC< TData > *)> fun)
- void set_kernel_fun (FUN< epiworld_double(VEC(epiworld_double)&, epiworld_double, LFMCMC< TData > *)> fun)
- const size t get n samples ()
- const size_t get_n_statistics ()
- const size_t get_n_parameters ()
- · const epiworld double get_epsilon ()
- const VEC (epiworld_double) &get_params_now()
- const VEC (epiworld double) &get params prev()
- const VEC (epiworld double) &get params init()
- const VEC (epiworld_double) &get_statistics_obs()
- · const VEC (epiworld double) &get statistics hist()
- · const VEC (bool) &get_statistics_accepted()
- const VEC (epiworld_double) &get_posterior_lf_prob()
- const VEC (epiworld double) &get acceptance prob()
- const VEC (epiworld_double) &get_drawn_prob()
- VEC (TData) *get sampled data()

Random number generation

Parameters

eng

- void set rand engine (std::mt19937 &eng)
- std::mt19937 * get_rand_endgine ()
- void seed (unsigned int s)
- void **set_rand_gamma** (epiworld_double alpha, epiworld_double beta)
- epiworld_double runif ()
- epiworld_double rnorm ()
- epiworld double **rnorm** (epiworld double mean, epiworld double sd)
- epiworld double rgamma ()
- epiworld_double rgamma (epiworld_double alpha, epiworld_double beta)

3.4.1 Detailed Description

template < typename TData > class LFMCMC < TData >

Likelihood-Free Markov Chain Monte Carlo.

Template Parameters

TData Type of data that is generated

The documentation for this class was generated from the following file:

• include/epiworld/math/lfmcmc.hpp

3.5 Location < TSeq > Class Template Reference

Public Member Functions

```
    add_person (Person< TSeq > &p)
    add_person (Person< TSeq > *p)
    size_t count () const
```

· void reset ()

The documentation for this class was generated from the following file:

• include/epiworld/location-bones.hpp

3.6 Model < TSeq > Class Template Reference

Core class of epiworld.

```
#include <model-bones.hpp>
```

Public Member Functions

```
    DataBase< TSeq > & get_db ()
```

- epiworld_double & **operator()** (std::string pname)
- size_t size () const
- void record_variant (Virus < TSeq > &v)
- void record_tool (Tool < TSeq > &t)
- int get_nvariants () const
- unsigned int get_ndays () const
- unsigned int **get_n_replicates** () const
- void set_ndays (unsigned int ndays)
- bool get_verbose () const
- void verbose_off ()
- void verbose_on ()
- int today () const

The current time of the model.

void write_data (std::string fn_variant_info, std::string fn_variant_hist, std::string fn_tool_info, std::string fn_tool_hist, std::string fn_total_hist, std::string fn_transmission, std::string fn_transition) const

```
Wrapper of DataBase::write_data
```

- std::map< std::string, epiworld_double > & params ()
- void reset ()

Reset the model.

- void print () const
- Model < TSeq > && clone () const

void **get_elapsed** (std::string unit="auto", epiworld_double *last_elapsed=nullptr, epiworld_double *total_
 elapsed=nullptr, std::string *unit_abbr=nullptr, bool print=true) const

- void add_global_action (std::function< void(Model< TSeq > *)> fun, int date)
 - Set a global action.
- void run_global_actions ()
- · void clear status set ()

Set the backup object

backup can be used to restore the entire object after a run. This can be useful if the user wishes to have individuals start with the same network from the beginning.

- void set backup ()
- · void restore backup ()

Random number generation

Parameters

| eng | Random number generator |
|-----|-------------------------|
| s | Seed |

- void set_rand_engine (std::mt19937 &eng)
- std::mt19937 * get_rand_endgine ()
- void **seed** (unsigned int s)
- void set_rand_gamma (epiworld_double alpha, epiworld_double beta)
- epiworld double runif ()
- epiworld_double rnorm ()
- epiworld_double rnorm (epiworld_double mean, epiworld_double sd)
- epiworld double rgamma ()
- epiworld_double **rgamma** (epiworld_double alpha, epiworld_double beta)

Add Virus/Tool to the model

This is done before the model has been initialized.

Parameters

| V | Virus to be added |
|--------|---|
| t | Tool to be added |
| preval | Initial prevalence (initial state.) It can be specified as a proportion (between zero and one,) or an integer indicating number of individuals. |

- void add_virus (Virus < TSeq > v, epiworld_double preval)
- void add_virus_n (Virus< TSeq > v, unsigned int preval)
- void add_tool (Tool < TSeq > t, epiworld_double preval)
- void add_tool_n (Tool< TSeq > t, unsigned int preval)

Accessing population of the model

Parameters

| fn | std::string Filename of the edgelist file. |
|------|--|
| skip | int Number of lines to skip in fn. |

Parameters

| directed | bool Whether the graph is directed or not. | |
|----------|---|--|
| min_id | int Minimum id number (if negative, the program will try to guess from the data.) | |
| max_id | _id int Maximum id number (if negative, the program will try to guess from the data.) | |
| al | AdjList to read into the model. | |

- void population_from_adjlist (std::string fn, int skip=0, bool directed=false, int min_id=-1, int max_id=-1)
- void population_from_adjlist (AdjList al)
- · bool is directed () const
- std::vector< Person< TSeq > > * get_population ()
- void population_smallworld (unsigned int n=1000, unsigned int k=5, bool d=false, epiworld_double p=. ← 01)

Functions to run the model

Parameters

| seed | Seed to be used for Pseudo-RNG. |
|-------|---|
| ndays | Number of days (steps) of the simulation. |
| fun | In the case of run_multiple, a function that is called after each experiment. |

- · void init (unsigned int ndays, unsigned int seed)
- void update_status ()
- void mutate_variant ()
- void next ()
- void run ()

Runs the simulation (after initialization)

 void run_multiple (unsigned int nexperiments, std::function< void(Model< TSeq > *)> fun, bool reset, bool verbose)

Rewire the network preserving the degree sequence.

This implementation assumes an undirected network, thus if $\{(i,j), (k,l)\} \rightarrow \{(i,l), (k,j)\}$, the reciprocal is also true, i.e., $\{(j,i), (l,k)\} \rightarrow \{(j,k), (l,i)\}$.

Parameters

| proportion | Proportion of ties to be rewired. |
|------------|-----------------------------------|

Returns

A rewired version of the network.

- void **set_rewire_fun** (std::function< void(std::vector< Person< TSeq >> *, Model< TSeq > *, epiworld double)> fun)
- void set_rewire_prop (epiworld_double prop)
- epiworld double get rewire prop () const
- · void rewire ()

Export the network data in edgelist form

Parameters

| | fn | std::string. File name. |
|--|----|-------------------------|
|--|----|-------------------------|

Parameters

| source | Integer vector |
|--------|----------------|
| target | Integer vector |

When passing the source and target, the function will write the edgelist on those.

- · void write edgelist (std::string fn) const
- void write_edgelist (std::vector< unsigned int > &source, std::vector< unsigned int > &target) const

Manage status (states) in the model

The functions get status return the current values for the statuses included in the model.

Parameters

```
lab std::string Name of the status.
```

Returns

```
add_status* returns nothing.
get_status_* returns a vector of pairs with the statuses and their labels.
```

- void add status (std::string lab, UpdateFun< TSeq > fun=nullptr)
- const std::vector< std::string > & get_status () const
- const std::vector< UpdateFun< TSeq > > & get_status_fun () const
- void print_status_codes () const

Set the user data object

Parameters

| names | string vector with the names of the variables. |
|-------|--|
|-------|--|

- void set_user_data (std::vector< std::string > names)
- void **add_user_data** (unsigned int j, epiworld_double x)
- void add_user_data (std::vector< epiworld_double > x)
- UserData < TSeq > & get_user_data ()

Queuing system

When queueing is on, the model will keep track of which agents are either in risk of exposure or exposed. This then is used at each step to act only on the aforementioned agents.

```
• void queuing_on ()
```

Activates the queuing system (default.)

void queuing_off ()

Deactivates the queuing system.

• bool is_queuing_on () const

Query if the queuing system is on.

Queue < TSeq > & get_queue ()

Retrieve the Queue object.

Get the susceptibility reduction object

Parameters



Returns

epiworld double

- void set_susceptibility_reduction_mixer (MixerFun< TSeq > fun)
- void set_transmission_reduction_mixer (MixerFun< TSeq > fun)
- void set_recovery_enhancer_mixer (MixerFun< TSeq > fun)
- void set_death_reduction_mixer (MixerFun< TSeq > fun)

Friends

- class Person < TSeq >
- class DataBase < TSeq >
- class Queue < TSeq >

Tool Mixers

These functions combine the effects tools have to deliver a single effect. For example, wearing a mask, been vaccinated, and the immune system combine together to jointly reduce the susceptibility for a given virus.

- std::vector< epiworld_double > array_double_tmp
- std::vector< VirusPtr< TSeq > * > array_virus_tmp
- · Model ()
- Model (const Model < TSeq > &m)
- Model (Model < TSeq > &&m)
- Model < TSeq > & operator= (const Model < TSeq > &m)
- void clone_population (std::vector< Person< TSeq > > &p, std::map< int, int > &p_ids, bool &d, Model<
 TSeq > *m=nullptr) const
- void clone_population (const Model < TSeq > &m)

Setting and accessing parameters from the model

Tools can incorporate parameters included in the model. Internally, parameters in the tool are stored as pointers to an std::map<> of parameters in the model. Using the unsigned int method directly fetches the parameters in the order these were added to the tool. Accessing parameters via the std::string method involves searching the parameter directly in the std::map<> member of the model (so it is not recommended.)

The function $set_param()$ can be used when the parameter already exists in the model.

The par() function members are aliases for get_param().

Parameters

| initial_val | |
|-------------|--|
| pname | Name of the parameter to add or to fetch |

Returns

The current value of the parameter in the model.

- epiworld_double * p0
- epiworld double * p1
- epiworld_double * p2
- epiworld double * p3
- epiworld_double * p4
- epiworld_double * p5
- epiworld double * p6
- epiworld double * p7
- epiworld double * p8
- epiworld_double * p9
- epiworld_double * p10
- epiworld double * p11
- epiworld_double * p12
- epiworld_double * p13
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- epiworld_double * p27
- epiworld_double * p28epiworld_double * p29
- epiworld_double * p30
- epiworld_double * p31
- epiworld double * p32
- epiworld double * p33
- epiworld double * p34
- epiworld_double * p35
- epiworld_double * p36
- epiworld double * p37
- epiworld double * p38
- epiworld double * p39
- unsigned int **npar_used** = 0u
- epiworld_double add_param (epiworld_double initial_val, std::string pname)
- epiworld_double set_param (std::string pname)
- epiworld double get_param (unsigned int k)
- epiworld_double get_param (std::string pname)
- epiworld_double **par** (unsigned int k)
- epiworld_double par (std::string pname)

3.6.1 Detailed Description

```
template<typename TSeq = bool> class Model< TSeq >
```

Core class of epiworld.

The model class provides the wrapper that puts together Person, Virus, and Tools.

Template Parameters

| TSeq | Type of sequence. In principle, users can build models in which virus and human sequence is |
|------|---|
| | represented as numeric vectors (if needed.) |

3.6.2 Member Function Documentation

3.6.2.1 add_global_action()

```
template<typename TSeq = bool>
void Model< TSeq >::add_global_action (
          std::function< void(Model< TSeq > *)> fun,
          int date )
```

Set a global action.

Parameters

| fun | A function to be called on the prescribed dates |
|------|--|
| date | Integer indicating when the function is called (see details) |

When date is less than zero, then the function is called at the end of every day. Otherwise, the function will be called only at the end of the indicated date.

3.6.2.2 reset()

```
template<typename TSeq = bool>
void Model< TSeq >::reset ( )
```

Reset the model.

Resetting the model will:

- · clear the database
- restore the population (if set_backup() was called before)

- · re-distribute tools
- · re-distribute viruses
- set the date to 0

3.6.2.3 run_multiple()

Parameters

| nexperiments | Multiple runs of the simulation | |
|--------------|---------------------------------|--|
|--------------|---------------------------------|--|

3.6.2.4 write_data()

Wrapper of DataBase::write_data

Parameters

| fn_variant_info | Filename. Information about the variant. |
|-----------------|--|
| fn_variant_hist | Filename. History of the variant. |
| fn_tool_info | Filename. Information about the tool. |
| fn_tool_hist | Filename. History of the tool. |
| fn_total_hist | Filename. Aggregated history (status) |
| fn_transmission | Filename. Transmission history. |
| fn_transition | Filename. Markov transition history. |

The documentation for this class was generated from the following files:

- include/epiworld/config.hpp
- include/epiworld/model-bones.hpp

3.7 Person < TSeq > Class Template Reference

Person (agents)

#include <person-bones.hpp>

Public Member Functions

- Person (const Person < TSeq > &p)
- int get_id () const

Id of the individual.

• unsigned int get_index () const

Location (0, ..., n-1).

- std::mt19937 * get_rand_endgine()
- Model < TSeq > * get_model ()
- VirusPtr< TSeq > & get_virus (int i)
- $std::vector < VirusPtr < TSeq > > & get_viruses ()$
- · size t get n viruses () const noexcept
- ToolPtr < TSeq > & get_tool (int i)
- std::vector< ToolPtr< TSeq > > & get_tools ()
- size t get n tools () const noexcept
- void mutate_variant ()
- void add neighbor (Person< TSeq > *p, bool check source=true, bool check target=true)
- std::vector< Person< TSeq > * > & get_neighbors ()
- void **change_status** (epiworld_fast_uint new_status, epiworld_fast_int queue=0)
- const epiworld_fast_uint & get_status () const
- · void reset ()
- · bool has tool (unsigned int t) const
- · bool has tool (std::string name) const
- · bool has virus (unsigned int t) const
- · bool has virus (std::string name) const
- · bool is_locked () const noexcept

Add/Remove Virus/Tool

Calling any of these functions will lock the agent (person) until the action is applied at the end of the iteration. Calling any of this functions when the agent is locked will cause an error.

Parameters

| tool | Tool to add |
|------------|-------------------------|
| virus | Virus to add |
| status_new | Status after the change |
| queue | |

- void **add_tool** (ToolPtr< TSeq > tool, epiworld_fast_int status_new=-99, epiworld_fast_int queue=-99)
- void add_tool (Tool < TSeq > tool, epiworld_fast_int status_new=-99, epiworld_fast_int gueue=-99)
- void add_virus (VirusPtr< TSeq > virus, epiworld_fast_int status_new=-99, epiworld_fast_int queue=-99)
- void add virus (Virus < TSeq > virus, epiworld fast int status new=-99, epiworld fast int queue=-99)
- void rm_tool (epiworld_fast_uint tool_idx, epiworld_fast_int status_new=-99, epiworld_fast_int queue=-99)

void rm_virus (epiworld_fast_uint virus_idx, epiworld_fast_int status_new=-99, epiworld_fast_int queue=-99)

Get the rates (multipliers) for the agent

Parameters

```
v A pointer to a virus.
```

Returns

epiworld_double

- epiworld_double **get_susceptibility_reduction** (VirusPtr< TSeq > v)
- epiworld_double get_transmission_reduction (VirusPtr < TSeq > v)
- epiworld_double get_recovery_enhancer (VirusPtr< TSeq > v)
- epiworld_double get_death_reduction (VirusPtr< TSeq > v)

Friends

- class Model < TSeq >
- class Tool < TSeq >
- class Queue < TSeq >
- void default_add_virus (Action < TSeq > &a, Model < TSeq > *m)
- void default_add_tool (Action < TSeq > &a, Model < TSeq > *m)
- void default_rm_virus (Action < TSeq > &a, Model < TSeq > *m)
- void default_rm_tool (Action < TSeq > &a, Model < TSeq > *m)

3.7.1 Detailed Description

```
template<typename TSeq = bool> class Person< TSeq >
```

Person (agents)

Template Parameters

```
TSeq | Sequence type (should match TSeq across the model)
```

The documentation for this class was generated from the following files:

- include/epiworld/config.hpp
- include/epiworld/person-bones.hpp

3.8 PersonTools < TSeq > Class Template Reference

List of tools available for the individual to.

#include <persontools-bones.hpp>

Public Member Functions

- void add_tool (int date, Tool < TSeq > tool)
- epiworld_double get_susceptibility_reduction (VirusPtr< TSeq > v)
- epiworld double get_transmission_reduction (VirusPtr< TSeq > v)
- epiworld_double get_recovery_enhancer (VirusPtr< TSeq > v)
- epiworld_double get_death_reduction (VirusPtr< TSeq > v)
- void set_susceptibility_reduction_mixer (MixerFun < TSeq > fun)
- void set_transmission_reduction_mixer (MixerFun< TSeq > fun)
- void set recovery enhancer mixer (MixerFun < TSeq > fun)
- void set_death_reduction_mixer (MixerFun < TSeq > fun)
- · size t size () const
- Tool < TSeq > & operator() (int i)
- Person < TSeq > * get_person ()
- Model < TSeq > * get_model ()
- void reset ()
- · bool has_tool (unsigned int t) const
- · bool has_tool (std::string name) const

Friends

- class Person < TSeq >
- class Model < TSeq >

3.8.1 Detailed Description

template < typename TSeq = bool > class PersonTools < TSeq >

List of tools available for the individual to.

Template Parameters

TSeq

The documentation for this class was generated from the following files:

- · include/epiworld/config.hpp
- · include/epiworld/persontools-bones.hpp
- include/epiworld/persontools-meat.hpp

3.9 PersonViruses < TSeq > Class Template Reference

Set of viruses in host.

#include <personviruses-bones.hpp>

Public Member Functions

- void add_virus (epiworld_fast_uint new_status, Virus < TSeq > v)
- size_t size () const
- int size active () const
- Virus< TSeq > & operator() (int i)
- · void mutate ()
- · void reset ()
- void deactivate (Virus < TSeq > &v)
- Person< TSeq > * get host ()
- bool has_virus (unsigned int v) const
- bool has_virus (std::string vname) const

Friends

- class Person < TSeq >
- class Model < TSeq >

3.9.1 Detailed Description

```
template<typename TSeq = bool> class PersonViruses< TSeq >
```

Set of viruses in host.

Template Parameters

```
TSeq Type of sequence
```

The documentation for this class was generated from the following files:

- include/epiworld/personviruses-bones.hpp
- include/epiworld/personviruses-meat.hpp

3.10 Progress Class Reference

A simple progress bar.

```
#include progress.hpp>
```

Public Member Functions

- Progress (int n_, int width_)
- void start ()
- void next ()
- void end ()

3.10.1 Detailed Description

A simple progress bar.

The documentation for this class was generated from the following file:

· include/epiworld/progress.hpp

3.11 Queue < TSeq > Class Template Reference

Controls which agents are verified at each step.

```
#include <queue-bones.hpp>
```

Public Member Functions

- void operator+= (Person< TSeq > *p)
- void operator-= (Person< TSeq > *p)
- epiworld fast int operator[] (unsigned int i) const
- void set_model (Model < TSeq > *m)

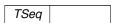
3.11.1 Detailed Description

```
template<typename TSeq = bool> class Queue< TSeq >
```

Controls which agents are verified at each step.

The idea is that only agents who are either in an infected state or have an infected neighbor should be checked. Otherwise it makes no sense (no chance to recover or capture the disease).

Template Parameters



The documentation for this class was generated from the following files:

- include/epiworld/model-bones.hpp
- · include/epiworld/queue-bones.hpp

3.12 RandGraph Class Reference

Public Member Functions

RandGraph (int N_)

- · void init (int s)
- void set_rand_engine (std::mt19937 &e)
- epiworld double runif ()

The documentation for this class was generated from the following file:

· include/epiworld/random graph.hpp

3.13 Tool < TSeq > Class Template Reference

Tools for defending the host against the virus.

```
#include <tools-bones.hpp>
```

Public Member Functions

- Tool (std::string name="unknown tool")
- void set_sequence (TSeq d)
- void set_sequence_unique (TSeq d)
- void set_sequence (std::shared_ptr< TSeq > d)
- std::shared_ptr< TSeq > get_sequence ()
- TSeq & get_sequence_unique ()
- void set_name (std::string name)
- std::string **get_name** () const
- Person < TSeq > * get_person ()
- void set_person (Person < TSeq > *p)
- unsigned int **get_id** () const
- void set_id (int id)
- · void set date (int d)
- · void set_status (epiworld_fast_int init, epiworld_fast_int post)
- void set_queue (epiworld_fast_int init, epiworld_fast_int post)
- void get_status (epiworld_fast_int *init, epiworld_fast_int *post)
- void **get_queue** (epiworld fast int *init, epiworld fast int *post)

Get and set the tool functions

Parameters

| ٧ | The virus over which to operate |
|-----|---------------------------------|
| fun | the function to be used |

Returns

epiworld double

- epiworld double get_susceptibility_reduction (VirusPtr< TSeq > v)
- epiworld_double get_transmission_reduction (VirusPtr< TSeq > v)
- epiworld_double get_recovery_enhancer (VirusPtr< TSeq > v)
- epiworld_double get_death_reduction (VirusPtr< TSeq > v)
- void set_susceptibility_reduction_fun (ToolFun < TSeq > fun)

- void set_transmission_reduction_fun (ToolFun< TSeq > fun)
- void set_recovery_enhancer_fun (ToolFun< TSeq > fun)
- void set_death_reduction_fun (ToolFun < TSeq > fun)
- void set_susceptibility_reduction (epiworld_double *prob)
- void set_transmission_reduction (epiworld_double *prob)
- void set_recovery_enhancer (epiworld_double *prob)
- void set_death_reduction (epiworld_double *prob)
- void set_susceptibility_reduction (epiworld_double prob)
- void set_transmission_reduction (epiworld_double prob)
- void set recovery enhancer (epiworld double prob)
- void set_death_reduction (epiworld_double prob)

Friends

- class PersonTools < TSeq >
- class Person < TSeq >
- class Model < TSeq >

3.13.1 Detailed Description

```
template < typename TSeq = bool > class Tool < TSeq >
```

Tools for defending the host against the virus.

Template Parameters

| TSeq | Type of sequence |
|------|------------------|
|------|------------------|

The documentation for this class was generated from the following files:

- include/epiworld/config.hpp
- include/epiworld/tools-bones.hpp
- include/epiworld/tools-meat.hpp

3.14 UserData < TSeq > Class Template Reference

Personalized data by the user.

```
#include <userdata-bones.hpp>
```

Public Member Functions

UserData (std::vector< std::string > names)

Construct a new User Data object.

- std::vector< std::string > & get_names ()
- std::vector< int > & get_dates ()
- std::vector< epiworld_double > & get_data ()

void **get_all** (std::vector< std::string > *names=nullptr, std::vector< int > *date=nullptr, std::vector<
 epiworld_double > *data=nullptr)

- unsigned int nrow () const
- unsigned int ncol () const
- void write (std::string fn)
- · void print () const

Append data

Parameters

| Х | A vector of length ncol() (if vector), otherwise a epiworld_double. |
|---|---|
| j | Index of the data point, from 0 to ncol () - 1. |

- void add (std::vector< epiworld_double > x)
- void **add** (unsigned int j, epiworld_double x)

Access data

Parameters

| i | Row (0 through ndays - 1.) | |
|---|-----------------------------|--|
| j | Column (0 through ncols()). | |

Returns

epiworld double&

- epiworld_double & operator() (unsigned int i, unsigned int j)
- epiworld double & operator() (unsigned int i, std::string name)

Friends

- class Model < TSeq >
- class ${\bf DataBase}{<}{\,{\sf TSeq}}{\,>}$

3.14.1 Detailed Description

template<typename TSeq> class UserData< TSeq>

Personalized data by the user.

Template Parameters

| TSeq | |
|------|--|

3.14.2 Constructor & Destructor Documentation

3.14.2.1 UserData()

Construct a new User Data object.

Parameters

names A vector of names. The length of the vector sets the number of columns to record.

The documentation for this class was generated from the following files:

- · include/epiworld/database-bones.hpp
- include/epiworld/userdata-bones.hpp
- include/epiworld/userdata-meat.hpp

3.15 vecHasher < T > Struct Template Reference

Vector hasher.

```
#include <misc.hpp>
```

Public Member Functions

• std::size_t operator() (std::vector< T > const &dat) const noexcept

3.15.1 Detailed Description

template<typename T> struct vecHasher< T>

Vector hasher.

Template Parameters



The documentation for this struct was generated from the following file:

include/epiworld/misc.hpp

3.16 Virus < TSeq > Class Template Reference

Virus.

```
#include <virus-bones.hpp>
```

Public Member Functions

- Virus (std::string name="unknown virus")
- · void mutate ()
- void set_mutation (MutFun < TSeq > fun)
- const TSeq * get_sequence ()
- void **set_sequence** (TSeq sequence)
- Person< TSeq > * get host ()
- void set_host (Person < TSeq > *p)
- Model < TSeq > * get_model ()
- void set_date (int d)
- int get_date () const
- void set_id (int idx)
- int get_id () const
- void set_name (std::string name)
- std::string get_name () const
- std::vector< epiworld_double > & get_data ()

Get and set the tool functions

Parameters

| V | The virus over which to operate | |
|-----|---------------------------------|--|
| fun | the function to be used | |

Returns

epiworld_double

- epiworld_double get_prob_infecting ()
- epiworld_double get_prob_recovery ()
- epiworld_double get_prob_death ()
- void post_recovery ()
- void set_post_recovery (PostRecoveryFun< TSeq > fun)
- void set_post_immunity (epiworld_double prob)
- void set_post_immunity (epiworld_double *prob)
- void set_prob_infecting_fun (VirusFun< TSeq > fun)
- void set_prob_recovery_fun (VirusFun < TSeq > fun)
- void set_prob_death_fun (VirusFun< TSeq > fun)
- void set_prob_infecting (epiworld_double *prob)
- void set_prob_recovery (epiworld_double *prob)
- void set_prob_death (epiworld_double *prob)
- void set_prob_infecting (epiworld_double prob)
- void set_prob_recovery (epiworld_double prob)
- void set_prob_death (epiworld_double prob)

Get and set the status and queue

After applied, viruses can change the status and affect the queue of agents. These function sets the default values, which are retrieved when adding or removing a virus does not specify a change in status or in queue.

Parameters

| init | After the virus/tool is added to the host. |
|---------|--|
| end | After the virus/tool is removed. |
| removed | After the host (Person) is removed. |

- void **set_status** (epiworld_fast_int init, epiworld_fast_int end, epiworld_fast_int removed)
- void **set_queue** (epiworld_fast_int init, epiworld_fast_int end, epiworld_fast_int removed)
- void get_status (epiworld_fast_int *init, epiworld_fast_int *end, epiworld_fast_int *removed)
- void get_queue (epiworld_fast_int *init, epiworld_fast_int *end, epiworld_fast_int *removed)

Friends

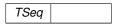
- class Person < TSeq >
- class Model < TSeq >
- class PersonViruses < TSeq >
- class DataBase< TSeq >

3.16.1 Detailed Description

template<typename TSeq = bool> class Virus< TSeq >

Virus.

Template Parameters



Raw transmisibility of a virus should be a function of its genetic sequence. Nonetheless, transmisibility can be reduced as a result of having one or more tools to fight the virus. Because of this, transmisibility should be a function of the host.

The documentation for this class was generated from the following files:

- include/epiworld/config.hpp
- include/epiworld/virus-bones.hpp
- · include/epiworld/virus-meat.hpp

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