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.pop_from_random(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:
Distribution of the population at time 100:
- Total healthy (S) : 99995 -> 97390
- Total recovered (S) : 0 -> 2554
- Total infected (I)
                    :
                            5 -> 56
 - Total removed (R)
                            0 -> 0
(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, Pv in [0,1],
- 2. The contagion reduction factor of i, Cr in [0,1], and
- 3. The host's transmission reduction factor, Tr [0,1].

The last two are computed from i and j's tools. Ultimately, the probability of i getting virus v from j equals: $P(Virus \ v) = Pv \ \star \ (1 - Cr) \ \star (1 - Tr)$

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:

```
P(Virus \ v | at most one virus) = Prcond(i, v, j)
```

```
The latter is calculated using Bayes' rule
```

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:

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PersonViruses < TSeq >	
Set of viruses in host	20
Progress	
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Queue < TSeq >	
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RandGraph	22
Tool< TSeq >	
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Virus< TSeq >	
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Chapter 3

Class Documentation

3.1 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

3.1.1 Constructor & Destructor Documentation

3.1.1.1 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

3.2 DataBase < TSeq > Class Template Reference

Statistical data about the process.

#include <database-bones.hpp>

Public Member Functions

- DataBase (int freq=1)
- void record_variant (Virus < TSeq > *v)

Registering a new variant.

- 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 up_exposed (Virus < TSeq > *v, epiworld_fast_uint new_status)
- void down_exposed (Virus< TSeq > *v, epiworld fast uint prev status)
- void state_change (epiworld_fast_uint prev_status, epiworld_fast_uint new_status)
- void record_transition (epiworld_fast_uint from, epiworld_fast_uint to)
- int get_today_total (std::string what) const

Get recorded information from the model.

- 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)
- void get_hist_variant (std::vector< int > &date, std::vector< int > &id, std::vector< std::string > &status, std::vector< int > &counts) const
- void write_data (std::string fn_variant_info, std::string fn_variant_hist, std::string fn_total_hist, std::string fn_transition)
 void write_data (std::string fn_variant_info, std::string fn_variant_hist, std::string fn_total_hist, std::string fn_transition)

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- 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 ()

Friends

class Model < TSeq >

3.2.1 Detailed Description

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

Statistical data about the process.

Template Parameters



3.2.2 Member Function Documentation

3.2.2.1 get_today_total()

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 @[
```

3.2.2.2 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

3.3 LFMCMC< TData > Class Template Reference

Likelihood-Free Markov Chain Monte Carlo.

```
#include <lfmcmc.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)
- void set_rand_engine (std::mt19937 &eng)

Random number generation.

- 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)
- const size_t get_n_samples ()

@1

- 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()

3.3.1 Detailed Description

```
template < typename TData > class LFMCMC < TData >
```

Likelihood-Free Markov Chain Monte Carlo.

Template Parameters

```
TData Type of data that is generated
```

3.3.2 Member Function Documentation

3.3.2.1 set_rand_engine()

Random number generation.

Parameters



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

• include/epiworld/math/lfmcmc.hpp

3.4 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

3.5 Model < TSeq > Class Template Reference

Public Member Functions

bool get_verbose () const

```
    Model (const Model < TSeg > &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 <</li>

 TSeq > *m=nullptr) const

    void clone_population (const Model < TSeq > &m)

· void set backup ()
     Set the backup object.

    void restore_backup ()

    DataBase< TSeq > & get_db ()

    epiworld double & operator() (std::string pname)

• size_t size () const
void set_rand_engine (std::mt19937 &eng)
     Random number generation.

    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)

    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)

    void pop_from_adjlist (std::string fn, int skip=0, bool directed=false, int min_id=-1, int max_id=-1)

     Accessing population of the model.

    void pop from adjlist (AdjList al)

· bool is_directed () const

    std::vector< Person< TSeq > > * get_population ()

    void pop_from_random (unsigned int n=1000, unsigned int k=5, bool d=false, epiworld_double p=.01)

• void init (unsigned int ndays, unsigned int seed)

    void update_status ()

    void mutate_variant()

· void next ()
· void run ()

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

    void record_variant (Virus< TSeq > *v)

• int get_nvariants () const
· unsigned int get ndays () const
· unsigned int get_n_replicates () const

    void set_ndays (unsigned int ndays)
```

- void verbose off ()
- void verbose_on ()
- int today () const
- void set_rewire_fun (std::function< void(std::vector< Person< TSeq >> *, Model< TSeq > *, epiworld_← double)> fun)

Rewire the network preserving the degree sequence.

- void set_rewire_prop (epiworld_double prop)
- epiworld_double get_rewire_prop () const
- · void rewire ()
- void set update susceptible (UpdateFun< TSeq > fun)

@

- void set_update_exposed (UpdateFun< TSeq > fun)
- void set update removed (UpdateFun< TSeq > fun)
- void write_data (std::string fn_variant_info, std::string fn_variant_hist, std::string fn_total_hist, std::string fn transmission, std::string fn transmission, std::string fn transmission

Wrapper of DataBase::write_data

• void write_edgelist (std::string fn) const

Export the network data in edgelist form.

- void write_edgelist (std::vector< unsigned int > &source, std::vector< unsigned int > &target) const
- std::map< std::string, epiworld_double > & params ()

@]

· void reset ()

Reset the model.

- · void print () const
- Model < TSeq > && clone () const
- void add_status_susceptible (epiworld_fast_uint s, std::string lab)

Adds extra statuses to the model.

- void add_status_exposed (epiworld_fast_uint s, std::string lab)
- void add_status_removed (epiworld_fast_uint s, std::string lab)
- void add_status_susceptible (std::string lab)
- void add_status_exposed (std::string lab)
- void add_status_removed (std::string lab)
- const std::vector< epiworld_fast_uint > & get_status_susceptible () const
- const std::vector< epiworld_fast_uint > & ${\tt get_status_exposed}$ () const
- const std::vector< epiworld_fast_uint > & **get_status_removed** () const
- const std::vector< std::string > & $get_status_susceptible_labels$ () const
- const std::vector< std::string > & get_status_exposed_labels () const
- const std::vector< std::string > & get_status_removed_labels () const
- · void print status codes () const
- epiworld_fast_uint get_default_susceptible () const
- epiworld_fast_uint get_default_exposed () const
- epiworld_fast_uint get_default_removed () const
- void reset_status_codes (std::vector< epiworld_fast_uint > codes, std::vector< std::string > names, bool verbose=true)

@]

• epiworld_double add_param (epiworld_double initial_val, std::string pname)

Setting and accessing parameters from the model.

- 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)

Public Attributes

bool is_queuing_on () constQueue< TSeq > & get_queue ()

```
    std::vector< epiworld_double > array_double_tmp

  std::vector < Virus < TSeq > * > array virus tmp
• 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
  epiworld_double * p14
• epiworld double * p15
• epiworld double * p16
  epiworld double * p17
• epiworld_double * p18
epiworld_double * p19
• epiworld_double * p20
• epiworld double * p21
• epiworld double * p22
• epiworld_double * p23
epiworld_double * p24
• epiworld_double * p25
• epiworld_double * p26
  epiworld double * p27
• epiworld_double * p28
• epiworld 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

Friends

```
class Person < TSeq >
```

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

3.5.1 Member Function Documentation

3.5.1.1 add_param()

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

initia	_val	
pnan	пе	Name of the parameter to add or to fetch

Returns

The current value of the parameter in the model. @[

3.5.1.2 add_status_susceptible()

Adds extra statuses to the model.

Adding values of ${\tt s}$ that are already present in the model will result in an error.

The functions get_status_* return the current values for the statuses included in the model.

Parameters

s	unsigned int Code of the status
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. @[
```

3.5.1.3 init()

```
template<typename TSeq = bool>
void Model< TSeq >::init (
          unsigned int ndays,
          unsigned int seed )
```

@]

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. @[

3.5.1.4 pop_from_adjlist()

```
int skip = 0,
bool directed = false,
int min_id = -1,
int max_id = -1)
```

Accessing population of the model.

Parameters

fn	std::string Filename of the edgelist file.	
skip	int Number of lines to skip in fn.	
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	int Maximum id number (if negative, the program will try to guess from the data.)	
al	AdjList to read into the model. @[

3.5.1.5 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

3.5.1.6 reset_status_codes()

@]

Reset all the status codes of the model

The default values are those specified in the enum STATUS.

Parameters

codes	In the following order: Susceptible, Infected, Removed
names	Names matching the codes
verbose	When true, it will print the new mappings.

3.5.1.7 set_backup()

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

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. @[

3.5.1.8 set_rand_engine()

```
template<typename TSeq = bool>
void Model< TSeq >::set_rand_engine (
          std::mt19937 & eng )
```

Random number generation.

Parameters



3.5.1.9 set_rewire_fun()

Rewire the network preserving the degree sequence.

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

Parameters

proportion	Proportion of ties to be rewired.
ριοροιτίστι	Froportion of ties to be rewired.

Returns

A rewired version of the network. @[

3.5.1.10 set_user_data()

Set the user data object.

Parameters

names [@

3.5.1.11 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_total_hist	Filename. Aggregated history (status)
fn_transmission	Filename. Transmission history.
fn_transition	Filename. Markov transition history.

3.5.1.12 write_edgelist()

Export the network data in edgelist form.

Parameters

fn	std::string. File name.
source	Integer vector
target	Integer vector

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

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

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

3.6 Person < TSeq > Class Template Reference

Public Member Functions

- void init (epiworld_fast_uint baseline_status)
- void add_tool (int d, Tool < TSeq > tool)
- void add_virus (Virus < TSeq > *virus)
- void rm_virus (Virus < TSeq > *virus)
- epiworld_double get_susceptibility_reduction (Virus< TSeq > *v)
- epiworld_double get_transmission_reduction (Virus< TSeq > *v)
- epiworld_double get_recovery_enhancer (Virus< TSeq > *v)
- epiworld_double get_death_reduction (Virus< TSeq > *v)
- int get_id () const
- unsigned int **get_index** () const
- std::mt19937 * get_rand_endgine ()
- Model < TSeq > * get_model ()
- Virus< TSeq > & get_virus (int i)
- $\bullet \ \ \mathsf{PersonViruses} < \mathsf{TSeq} > \& \ \mathbf{get_viruses} \ ()$
- Tool < TSeq > & get_tool (int i)
- PersonTools< TSeq > & get_tools ()
- void mutate_variant ()
- void add neighbor (Person < TSeq > *p, bool check source=true, bool check target=true)
- std::vector< Person< $TSeq > * > & get_neighbors ()$
- void update_status ()
- void update_status (epiworld_fast_uint new_status)
- const epiworld_fast_uint & get_status () const
- const epiworld_fast_uint & get_status_next () const
- void reset ()
- void set_update_susceptible (UpdateFun < TSeq > fun)
- void set_update_exposed (UpdateFun< TSeq > fun)
- void set_update_removed (UpdateFun< TSeq > fun)
- · 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 visited () const
- void toggle_visited ()

Friends

- class Model < TSeq >
- class Tool < TSeq >
- class Queue < TSeq >

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

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

3.7 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 (Virus < TSeq > *v)
- epiworld_double get_transmission_reduction (Virus< TSeq > *v)
- epiworld_double get_recovery_enhancer (Virus< TSeq > *v)
- epiworld double get_death_reduction (Virus < 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 >

3.7.1 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

3.8 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 >

3.8.1 Detailed Description

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

Set of viruses in host.

Template Parameters



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

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

3.9 Progress Class Reference

A simple progress bar.

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

Public Member Functions

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

3.9.1 Detailed Description

A simple progress bar.

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

· include/epiworld/progress.hpp

3.10 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)
- · void update ()

3.10.1 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

TSea	
1009	

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

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

3.11 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

3.12 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 ()
- epiworld_double get_susceptibility_reduction (Virus < TSeq > *v)

Get and set the tool functions.

- epiworld_double $\mbox{get_transmission_reduction} \ (\mbox{Virus} < \mbox{TSeq} > * \mbox{v})$
- epiworld_double get_recovery_enhancer (Virus < TSeq > *v)
- epiworld_double get_death_reduction (Virus < 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)
- void set_name (std::string name)

@]

- std::string get_name () const
- Person< TSeq > * get_person ()
- unsigned int get_id () const

Friends

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

3.12.1 Detailed Description

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

Tools for defending the host against the virus.

Template Parameters

```
TSeq Type of sequence
```

3.12.2 Member Function Documentation

3.12.2.1 get_susceptibility_reduction()

Get and set the tool functions.

Parameters

ν	,	The virus over which to operate
f	un	the function to be used

Returns

```
epiworld_double @[
```

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

3.13 UserData < TSeq > Class Template Reference

Public Member Functions

- UserData (std::vector< std::string > names)
- void add (std::vector< epiworld_double > x)
- void **add** (unsigned int j, epiworld double x)
- epiworld_double & operator() (unsigned int i, unsigned int j)
- epiworld_double & operator() (unsigned int i, std::string name)
- 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

Friends

- class Model < TSeq >
- class DataBase < TSeq >

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

3.14 vecHasher< T > Struct Template Reference

Public Member Functions

std::size t operator() (std::vector< T > const &dat) const noexcept

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

include/epiworld/misc.hpp

3.15 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 ()
- Model < TSeq > * get_model ()
- void set date (int d)
- int get_date () const
- void set_id (int idx)
- int get_id () const
- bool is_active () const
- void deactivate ()
- epiworld_double get_prob_infecting ()

Get and set the tool functions.

- 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 ${\bf set_prob_death_fun}$ (VirusFun< ${\sf TSeq} > {\sf fun}$)
- 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)
- void set_name (std::string name)

@i

- std::string **get_name** () const
- std::vector< epiworld_double > & get_data ()

Friends

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

3.15.1 Detailed Description

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

Virus.

Template Parameters

TCoa	
1364	

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.

3.15.2 Member Function Documentation

3.15.2.1 get_prob_infecting()

```
template<typename TSeq >
epiworld_double Virus< TSeq >::get_prob_infecting [inline]
```

Get and set the tool functions.

Parameters

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

Returns

epiworld_double @[

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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