barry: Your go-to motif accountant 0.0-1

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Main Page

Barry: your to-go motif accountant

This repository contains a C++ template library that essentially counts sufficient statistics on binary arrays. The idea of the library is that this can be used together to build exponential family models as those in Exponential Random Graph Models (ERGMs), but as a generalization that also deals with non square arrays.

Examples

Counting statistics in a graph

In the following code we create an array of size 5x5 of class Network (available in the namespace netcounters), add/remove ties, print the graph, and count common statistics used in ERGMs:

```
#include <iostream>
#include <ostream>
#include "../include/barry.hpp"
typedef std::vector< unsigned int > vuint;
int main() {
  // Creating network of size six with five ties
  netcounters::Network net(
       6, 6,
       {0, 0, 4, 4, 2, 0, 1},
{1, 2, 0, 2, 4, 0, 1}
  // How does this looks like?
std::cout « "Current view" « std::endl;
  net.print();
  // Adding extra ties
  net += {1, 0};
net(2, 0) = true;
  // And removing a couple
  net(0, 0) = false;
net -= {1, 1};
  std::cout « "New view" « std::endl;
  net.print();
  \ensuremath{//} Initializing the data. The program deals with freing the memory
  net.set_data(new netcounters::NetworkData, true);
  // Creating counter object for the network and adding stats to count netcounters::NetStatsCounter counter(&net);
  netcounters::counter_edges(counter.counters);
```

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```
netcounters::counter_ttriads(counter.counters);
  netcounters::counter_isolates(counter.counters);
  netcounters::counter_ctriads(counter.counters);
  netcounters::counter_mutual(counter.counters);
  // Counting and printing the results
  std::vector< double > counts = counter.count_all();
  std::cout «
    "Edges : " « counts[0] « std::endl «
"Transitive triads : " « counts[1] « std::endl «
"Isolates : " « counts[2] « std::endl «
"C triads : " « counts[3] « std::endl «
"Mutuals : " « counts[4] « std::endl;
 return 0;
Compiling this program using g++
g++ -std=c++11 -Wall -pedantic 08-counts.cpp -o counts && ./counts
Yields the following output:
Current view
                1
  0,] 1 1
   1,] .
            1
  . 1
   5,]
New view
  0,] .
1,] 1
            1 1
  1,] 1 .
2,] 1 .
                    . 1
   3,] .
4,] 1
           . 1 . .
   5,] . . . .
Edges
Transitive triads : 3
Isolates
C triads
```

Code of Conduct

Mutuals

Please note that the barry project is released with a Contributor Code of Conduct. By contributing to this project, you agree to abide by its terms.

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2.1 Modules

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4.1 File List

Here is a list of all files with brief descriptions:

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include/barry/model-meat.hpp	
include/barry/powerset-bones.hpp	
include/barry/powerset-meat.hpp	
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Module Documentation

5.1 Counting

Classes

- · class NetworkData
 - Data class for Networks.
- · class NodeData

Data definition for the PhyloArray class.

class Counter< Array_Type, Data_Type >

A counter function based on change statistics.

5.1.1 Detailed Description

barry includes a flexible way to generate counters based on change statistics. Since most of the time we are counting many motifs in a graph, change statistics make a reasonable (and efficient) way to make such counts.

In particular, let the motif be defined as s(y), with y as the binary array. The change statistic when adding cell y_{ij} , i.e. when the cell moves from being emty to have a one, is defined as

$$\delta(y_{ij}) = s_{ij}^{+}(y) - s_{ij}^{-}(y),$$

where $s^+_{ij}(y)$ and $s^-_{ij}(y)$ represent the motif statistic with and without the ij-cell. For example, in the case of networks, the change statistic for the number of edges is always 1.

To count statistics in an array, the [Counter] class will empty the array, initialize the counters, and then start counting while adding at each step a single cell, until matching the original array.

5.2 Statistical Models

Statistical models available in barry.

Classes

class Model < Array_Type, Data_Counter_Type, Data_Rule_Type >

General framework for discrete exponential models. This class allows generating discrete exponential models in the form of a linear exponential model:

class Flock

A Flock is a group of Geese.

· class Geese

Annotated Phylo Model.

5.2.1 Detailed Description

Statistical models available in barry.

5.3 Network counters

Counters for network models.

Functions

• void counter_edges (NetCounters *counters)

Number of edges.

void counter isolates (NetCounters *counters)

Number of isolated vertices.

void counter_mutual (NetCounters *counters)

Number of mutual ties.

- void counter_istar2 (NetCounters *counters)
- void counter_ostar2 (NetCounters *counters)
- void counter_ttriads (NetCounters *counters)
- void counter_ctriads (NetCounters *counters)
- void counter_density (NetCounters *counters)
- void counter_idegree15 (NetCounters *counters)
- void counter_odegree15 (NetCounters *counters)
- void counter_absdiff (NetCounters *counters, uint attr_id, double alpha=1.0)

Sum of absolute attribute difference between ego and alter.

• void counter_diff (NetCounters *counters, uint attr_id, double alpha=1.0, double tail_head=true)

Sum of attribute difference between ego and alter to pow(alpha)

- NETWORK_COUNTER (init_single_attr)
- void counter nodeicov (NetCounters *counters, uint attr id)
- void counter_nodeocov (NetCounters *counters, uint attr_id)
- void counter_nodecov (NetCounters *counters, uint attr_id)
- void counter_nodematch (NetCounters *counters, uint attr_id)
- void counter_idegree (NetCounters *counters, std::vector< uint > d)

Counts number of vertices with a given in-degree.

void counter_odegree (NetCounters *counters, std::vector< uint > d)

Counts number of vertices with a given out-degree.

void counter degree (NetCounters *counters, std::vector< uint > d)

Counts number of vertices with a given out-degree.

5.3.1 Detailed Description

Counters for network models.

5.3 Network counters

Parameters

counters | A pointer to a NetCounters object (Counters<Network, NetCounterData>).

5.3.2 Function Documentation

5.3.2.1 counter_absdiff()

Sum of absolute attribute difference between ego and alter.

Definition at line 419 of file network.hpp.

5.3.2.2 counter_ctriads()

Definition at line 322 of file network.hpp.

5.3.2.3 counter_degree()

Counts number of vertices with a given out-degree.

Definition at line 690 of file network.hpp.

5.3.2.4 counter_density()

Definition at line 361 of file network.hpp.

5.3.2.5 counter_diff()

Sum of attribute difference between ego and alter to pow(alpha)

Definition at line 461 of file network.hpp.

5.3.2.6 counter_edges()

Number of edges.

Definition at line 128 of file network.hpp.

5.3.2.7 counter_idegree()

Counts number of vertices with a given in-degree.

Definition at line 604 of file network.hpp.

5.3.2.8 counter_idegree15()

Definition at line 377 of file network.hpp.

5.3.2.9 counter_isolates()

Number of isolated vertices.

Definition at line 142 of file network.hpp.

5.3 Network counters

5.3.2.10 counter_istar2()

Definition at line 210 of file network.hpp.

5.3.2.11 counter_mutual()

Number of mutual ties.

Definition at line 172 of file network.hpp.

5.3.2.12 counter_nodecov()

Definition at line 558 of file network.hpp.

5.3.2.13 counter_nodeicov()

Definition at line 520 of file network.hpp.

5.3.2.14 counter_nodematch()

Definition at line 578 of file network.hpp.

5.3.2.15 counter_nodeocov()

Definition at line 539 of file network.hpp.

5.3.2.16 counter_odegree()

Counts number of vertices with a given out-degree.

Definition at line 646 of file network.hpp.

5.3.2.17 counter_odegree15()

Definition at line 397 of file network.hpp.

5.3.2.18 counter_ostar2()

Definition at line 228 of file network.hpp.

5.3.2.19 counter_ttriads()

Definition at line 247 of file network.hpp.

5.4 Phylo counters 15

5.3.2.20 NETWORK_COUNTER()

Definition at line 503 of file network.hpp.

5.4 Phylo counters

Counters for phylogenetic modeling.

Functions

- void counter_overall_gains (PhyloCounters *counters, bool duplication=true)

 Overall functional gains.
- void counter_gains (PhyloCounters *counters, std::vector< uint > nfun, bool duplication=true)

 Functional gains for a specific function (nfun).
- void counter_gains_k_offspring (PhyloCounters *counters, std::vector< uint > nfun, uint k=1u, bool duplication=true)

Functional gains for a specific function (nfun).

void counter_genes_changing (PhyloCounters *counters, bool duplication=true)

Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)

void counter_overall_loss (PhyloCounters *counters, bool duplication=true)

Overall functional loss.

- void counter_maxfuns (PhyloCounters *counters, uint lb, uint ub, bool duplication=true)
 - Cap the number of functions per gene.
- void counter loss (PhyloCounters *counters, std::vector< uint > nfun, bool duplication=true)

Total count of losses for an specific function.

void counter_overall_changes (PhyloCounters *counters, bool duplication=true)

Total number of changes. Use this statistic to account for "preservation".

- void counter_subfun (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)
 - Total count of Sub-functionalization events.
- void counter_cogain (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)

Co-evolution (joint gain or loss)

void counter_longest (PhyloCounters *counters)

Longest branch mutates (either by gain or by loss)

• void counter_neofun (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)

Total number of neofunctionalization events.

• void counter_neofun_a2b (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)

Total number of neofunctionalization events.

void counter_co_opt (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)
 Function co-opting.

5.4.1 Detailed Description

Counters for phylogenetic modeling.

Parameters

counters A pointer to a PhyloCounters object (Counters < PhyloArray, PhyloCounterData >).

5.4.2 Function Documentation

5.4.2.1 counter_co_opt()

Function co-opting.

Function co-opting of functions A and B happens when, for example, function B is gained as a new featured leveraging what function A already does; without losing function A. The sufficient statistic is defined as follows:

$$x_{pa}(1 - x_{pb}) \sum_{i < j} \left[x_{ia}^p (1 - x_{ib}^p) x_{ja}^p x_{jb}^p + x_{ja}^p (1 - x_{jb}^p) x_{ia}^p x_{ib}^p \right]$$

This algorithm implements the change statistic.

Definition at line 939 of file phylo.hpp.

5.4.2.2 counter_cogain()

Co-evolution (joint gain or loss)

Needs to specify pairs of functions (nfunA, nfunB).

Definition at line 626 of file phylo.hpp.

5.4 Phylo counters

5.4.2.3 counter_gains()

Functional gains for a specific function (nfun).

Definition at line 140 of file phylo.hpp.

5.4.2.4 counter_gains_k_offspring()

Functional gains for a specific function (nfun).

Definition at line 175 of file phylo.hpp.

5.4.2.5 counter_genes_changing()

Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)

Definition at line 238 of file phylo.hpp.

5.4.2.6 counter_longest()

Longest branch mutates (either by gain or by loss)

Definition at line 672 of file phylo.hpp.

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5.4.2.7 counter_loss()

Total count of losses for an specific function.

Definition at line 452 of file phylo.hpp.

5.4.2.8 counter_maxfuns()

Cap the number of functions per gene.

Definition at line 379 of file phylo.hpp.

5.4.2.9 counter_neofun()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 758 of file phylo.hpp.

5.4.2.10 counter_neofun_a2b()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 831 of file phylo.hpp.

5.4 Phylo counters

5.4.2.11 counter_overall_changes()

Total number of changes. Use this statistic to account for "preservation".

Definition at line 494 of file phylo.hpp.

5.4.2.12 counter_overall_gains()

Overall functional gains.

Total number of gains (irrespective of the function).

Definition at line 107 of file phylo.hpp.

5.4.2.13 counter_overall_loss()

Overall functional loss.

Definition at line 338 of file phylo.hpp.

5.4.2.14 counter_subfun()

Total count of Sub-functionalization events.

It requires to specify data = {funA, funB}

Definition at line 553 of file phylo.hpp.

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Chapter 6

Namespace Documentation

6.1 barry Namespace Reference

barry: Your go-to motif accountant

Namespaces

counters

Tree class and Treelterator class.

6.1.1 Detailed Description

barry: Your go-to motif accountant

6.2 barry::counters Namespace Reference

Tree class and Treelterator class.

Namespaces

- network
- phylo

6.2.1 Detailed Description

Tree class and Treelterator class.

6.3 barry::counters::network Namespace Reference

6.4 barry::counters::phylo Namespace Reference

6.5 CHECK Namespace Reference

Integer constants used to specify which cell should be check.

Variables

- const int BOTH = -1
- const int NONE = 0
- const int ONE = 1
- const int TWO = 2

6.5.1 Detailed Description

Integer constants used to specify which cell should be check.

6.5.2 Variable Documentation

6.5.2.1 BOTH

```
const int CHECK::BOTH = -1
```

Definition at line 31 of file typedefs.hpp.

6.5.2.2 NONE

```
const int CHECK::NONE = 0
```

Definition at line 32 of file typedefs.hpp.

6.5.2.3 ONE

```
const int CHECK::ONE = 1
```

Definition at line 33 of file typedefs.hpp.

6.5.2.4 TWO

```
const int CHECK::TWO = 2
```

Definition at line 34 of file typedefs.hpp.

6.6 EXISTS Namespace Reference

Integer constants used to specify which cell should be check to exist or not.

Variables

```
• const int BOTH = -1
```

- const int NONE = 0
- const int ONE = 1
- const int TWO = 1
- const int UKNOWN = -1
- const int AS_ZERO = 0
- const int AS_ONE = 1

6.6.1 Detailed Description

Integer constants used to specify which cell should be check to exist or not.

6.6.2 Variable Documentation

6.6.2.1 AS_ONE

```
const int EXISTS::AS_ONE = 1
```

Definition at line 49 of file typedefs.hpp.

6.6.2.2 AS_ZERO

```
const int EXISTS::AS_ZERO = 0
```

Definition at line 48 of file typedefs.hpp.

6.6.2.3 BOTH

```
const int EXISTS::BOTH = -1
```

Definition at line 42 of file typedefs.hpp.

6.6.2.4 NONE

```
const int EXISTS::NONE = 0
```

Definition at line 43 of file typedefs.hpp.

6.6.2.5 ONE

```
const int EXISTS::ONE = 1
```

Definition at line 44 of file typedefs.hpp.

6.6.2.6 TWO

```
const int EXISTS::TWO = 1
```

Definition at line 45 of file typedefs.hpp.

6.6.2.7 UKNOWN

```
const int EXISTS::UKNOWN = -1
```

Definition at line 47 of file typedefs.hpp.

Chapter 7

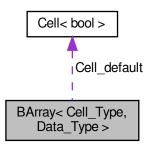
Class Documentation

7.1 BArray < Cell_Type, Data_Type > Class Template Reference

Baseline class for binary arrays.

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

 ${\tt Collaboration\ diagram\ for\ BArray}{<\tt Cell_Type,\ Data_Type>:}$



Public Member Functions

- bool operator== (const BArray< Cell_Type, Data_Type > &Array_)
- ∼BArray ()
- void set_data (Data_Type *data_, bool delete_data_=false)
- void out_of_range (uint i, uint j) const
- Cell_Type get_cell (uint i, uint j, bool check_bounds=true) const
- const Row_type< Cell_Type > * get_row (uint i, bool check_bounds=true) const
- const Col_type< Cell_Type > * get_col (uint i, bool check_bounds=true) const
- std::vector< Cell_Type > get_col_vec (uint i, bool check_bounds=true) const
- std::vector< Cell_Type > get_row_vec (uint i, bool check_bounds=true) const
- void get_col_vec (std::vector< Cell_Type > *x, uint i, bool check_bounds=true) const

- void get_row_vec (std::vector < Cell_Type > *x, uint i, bool check_bounds=true) const
- const Row_type< Cell_Type > & row (uint i, bool check_bounds=true) const
- const Col type < Cell Type > & col (uint i, bool check bounds=true) const
- Entries < Cell_Type > get_entries () const

Get the edgelist.

- · void transpose ()
- void clear (bool hard=true)
- void resize (uint N , uint M)
- void reserve ()
- void print () const

Constructors

Parameters

N_	Number of rows
M_	Number of columns
source	An unsigned vector ranging from 0 to N_
target	An unsigned int vector ranging from 0 to M_
target	When true tries to add repeated observations.

• BArray ()

Zero-size array.

BArray (uint N_, uint M_)

Empty array.

• BArray (uint N_, uint M_, const std::vector< uint > &source, const std::vector< uint > &target, const std::vector< Cell_Type > &value, bool add=true)

Edgelist with data.

BArray (uint N_, uint M_, const std::vector< uint > &source, const std::vector< uint > &target, bool add=true)

Edgelist with no data (simpler)

- BArray (const BArray < Cell_Type, Data_Type > &Array_, bool copy_data=false)
 Copy constructor.
- BArray< Cell_Type, Data_Type > & operator= (const BArray< Cell_Type, Data_Type > &Array_)
 Assignment constructor.

Move operator.

BArray< Cell_Type, Data_Type > & operator= (BArray< Cell_Type, Data_Type > &&x) noexcept
 Move assignment.

Queries

is_empty queries a single cell. nrow, ncol, and nnozero return the number of rows, columns, and non-zero cells respectively.

Parameters

i,j	Coordinates
check bounds	If false avoids checking bounds.

- bool is_empty (uint i, uint j, bool check_bounds=true) const
- · uint nrow () const noexcept
- uint ncol () const noexcept
- uint nnozero () const noexcept

Cell-wise insertion/deletion

Parameters

i,j	Row,column	
check_bounds	When true and out of range, the function throws an error.	
check_exists	Wither check if the cell exists (before trying to delete/add), or, in the case of swap_cells, check if either of both cells exists/don't exist.	

- BArray< Cell_Type, Data_Type > & operator+= (const std::pair< uint, uint > &coords)
- BArray< Cell_Type, Data_Type > & operator-= (const std::pair< uint, uint > &coords)
- BArrayCell< Cell Type, Data Type > operator() (uint i, uint j, bool check bounds=true)
- const BArrayCell const< Cell Type, Data Type > operator() (uint i, uint j, bool check bounds=true) const
- void rm_cell (uint i, uint j, bool check_bounds=true, bool check_exists=true)
- void insert_cell (uint i, uint j, const Cell< Cell_Type > &v, bool check_bounds, bool check_exists)
- void insert_cell (uint i, uint j, Cell< Cell_Type > &&v, bool check_bounds, bool check_exists)
- void insert_cell (uint i, uint j, Cell_Type v, bool check_bounds, bool check_exists)
- void swap_cells (uint i0, uint j0, uint i1, uint j1, bool check_bounds=true, int check_exists=CHECK::BOTH, int *report=nullptr)
- void toggle cell (uint i, uint j, bool check bounds=true, int check exists=EXISTS::UKNOWN)
- void toggle lock (uint i, uint j, bool check bounds=true)

Column/row wise interchange

- void swap_rows (uint i0, uint i1, bool check_bounds=true)
- void swap_cols (uint j0, uint j1, bool check_bounds=true)
- void zero row (uint i, bool check bounds=true)
- void zero_col (uint j, bool check_bounds=true)

Arithmetic operators

- BArray < Cell_Type, Data_Type > & operator+= (const BArray < Cell_Type, Data_Type > &rhs)
- BArray< Cell_Type, Data_Type > & operator+= (const Cell_Type &rhs)
- BArray< Cell_Type, Data_Type > & operator-= (const BArray< Cell_Type, Data_Type > &rhs)
- BArray< Cell_Type, Data_Type > & operator-= (const Cell_Type &rhs)
- BArray< Cell_Type, Data_Type > & operator/= (const Cell_Type &rhs)
- BArray< Cell_Type, Data_Type > & operator*= (const Cell_Type &rhs)

Public Attributes

- uint N
- uint M
- uint NCells = 0u
- std::vector< Row_type< Cell_Type > > el_ij
- std::vector< Col_type< Cell_Type > > el_ji
- Data_Type * data = nullptr
- bool delete_data = false
- bool visited = false

Static Public Attributes

static Cell
 Cell_Type > Cell_default

Friends

- class BArrayCell
 Cell_Type, Data_Type
- class BArrayCell_const< Cell_Type, Data_Type >

7.1.1 Detailed Description

```
template<typename Cell_Type = bool, typename Data_Type = bool> class BArray< Cell_Type, Data_Type >
```

Baseline class for binary arrays.

BArray class objects are arbitrary arrays in which non-empty cells hold data of type $Cell_Type$. The non-empty cells are stored by row and indexed using unordered_maps, i.e. $std::vector < std::unordered_{\leftarrow} map < unsigned int, <math>Cell_Type > .$

Template Parameters

Cell_Type	Type of cell (any type).
Data_Type	Data type of the array (bool default).

Definition at line 22 of file barray-bones.hpp.

7.1.2 Constructor & Destructor Documentation

7.1.2.1 BArray() [1/6]

```
template<typename Cell_Type = bool, typename Data_Type = bool>
BArray< Cell_Type, Data_Type >::BArray ( ) [inline]
```

Zero-size array.

Definition at line 58 of file barray-bones.hpp.

7.1.2.2 BArray() [2/6]

Empty array.

Definition at line 61 of file barray-bones.hpp.

7.1.2.3 BArray() [3/6]

Edgelist with data.

7.1.2.4 BArray() [4/6]

Edgelist with no data (simpler)

7.1.2.5 BArray() [5/6]

Copy constructor.

7.1.2.6 BArray() [6/6]

Move operator.

7.1.2.7 ∼BArray()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
BArray< Cell_Type, Data_Type >::~BArray ( )
```

7.1.3 Member Function Documentation

7.1.3.1 clear()

7.1.3.2 col()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
const Col_type< Cell_Type >& BArray< Cell_Type, Data_Type >::col (
    uint i,
    bool check_bounds = true) const
```

7.1.3.3 get_cell()

7.1.3.4 get_col()

7.1.3.5 get_col_vec() [1/2]

7.1.3.6 get_col_vec() [2/2]

7.1.3.7 get_entries()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Entries<Cell_Type> BArray< Cell_Type, Data_Type >::get_entries ( ) const
```

Get the edgelist.

Entries is a class with three objects: Two std::vector with the row and column coordinates respectively, and one std::vector with the corresponding value of the cell.

Returns

Entries<Cell_Type>

7.1.3.8 get_row()

7.1.3.9 get_row_vec() [1/2]

7.1.3.10 get_row_vec() [2/2]

7.1.3.11 insert_cell() [1/3]

7.1.3.12 insert_cell() [2/3]

7.1.3.13 insert_cell() [3/3]

7.1.3.14 is_empty()

7.1.3.15 ncol()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
uint BArray< Cell_Type, Data_Type >::ncol ( ) const [noexcept]
```

7.1.3.16 nnozero()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
uint BArray< Cell_Type, Data_Type >::nnozero ( ) const [noexcept]
```

7.1.3.17 nrow()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
uint BArray< Cell_Type, Data_Type >::nrow ( ) const [noexcept]
```

7.1.3.18 operator()() [1/2]

7.1.3.19 operator()() [2/2]

7.1.3.20 operator*=()

7.1.3.21 operator+=() [1/3]

7.1.3.22 operator+=() [2/3]

7.1.3.23 operator+=() [3/3]

7.1.3.24 operator-=() [1/3]

7.1.3.25 operator-=() [2/3]

7.1.3.26 operator-=() [3/3]

7.1.3.27 operator/=()

7.1.3.28 operator=() [1/2]

Move assignment.

7.1.3.29 operator=() [2/2]

Assignment constructor.

7.1.3.30 operator==()

7.1.3.31 out_of_range()

7.1.3.32 print()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
void BArray< Cell_Type, Data_Type >::print ( ) const
```

7.1.3.33 reserve()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
void BArray< Cell_Type, Data_Type >::reserve ( )
```

7.1.3.34 resize()

7.1.3.35 rm_cell()

7.1.3.36 row()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
const Row_type< Cell_Type >& BArray< Cell_Type, Data_Type >::row (
    uint i,
    bool check_bounds = true ) const
```

7.1.3.37 set_data()

7.1.3.38 swap_cells()

7.1.3.39 swap_cols()

7.1.3.40 swap_rows()

7.1.3.41 toggle_cell()

7.1.3.42 toggle_lock()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
void BArray< Cell_Type, Data_Type >::toggle_lock (
         uint i,
         uint j,
         bool check_bounds = true )
```

7.1.3.43 transpose()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
void BArray< Cell_Type, Data_Type >::transpose ( )
```

7.1.3.44 zero col()

7.1.3.45 zero_row()

7.1.4 Friends And Related Function Documentation

7.1.4.1 BArrayCell < Cell_Type, Data_Type >

```
template<typename Cell_Type = bool, typename Data_Type = bool>
friend class BArrayCell< Cell_Type, Data_Type > [friend]
```

Definition at line 1 of file barray-bones.hpp.

7.1.4.2 BArrayCell_const < Cell_Type, Data_Type >

```
template<typename Cell_Type = bool, typename Data_Type = bool>
friend class BArrayCell_const< Cell_Type, Data_Type > [friend]
```

Definition at line 1 of file barray-bones.hpp.

7.1.5 Member Data Documentation

7.1.5.1 Cell_default

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Cell< Cell_Type > BArray< Cell_Type, Data_Type >::Cell_default [static]
```

Definition at line 34 of file barray-bones.hpp.

7.1.5.2 data

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Data_Type* BArray< Cell_Type, Data_Type >::data = nullptr
```

Definition at line 31 of file barray-bones.hpp.

7.1.5.3 delete data

```
template<typename Cell_Type = bool, typename Data_Type = bool>
bool BArray< Cell_Type, Data_Type >::delete_data = false
```

Definition at line 32 of file barray-bones.hpp.

7.1.5.4 el_ij

```
template<typename Cell_Type = bool, typename Data_Type = bool>
std::vector< Row_type< Cell_Type > > BArray< Cell_Type, Data_Type >::el_ij
```

Definition at line 29 of file barray-bones.hpp.

7.1.5.5 el_ji

```
template<typename Cell_Type = bool, typename Data_Type = bool>
std::vector< Col_type< Cell_Type > > BArray< Cell_Type, Data_Type >::el_ji
```

Definition at line 30 of file barray-bones.hpp.

7.1.5.6 M

```
template<typename Cell_Type = bool, typename Data_Type = bool>
uint BArray< Cell_Type, Data_Type >::M
```

Definition at line 27 of file barray-bones.hpp.

7.1.5.7 N

```
template<typename Cell_Type = bool, typename Data_Type = bool>
uint BArray< Cell_Type, Data_Type >::N
```

Definition at line 26 of file barray-bones.hpp.

7.1.5.8 NCells

```
template<typename Cell_Type = bool, typename Data_Type = bool>
uint BArray< Cell_Type, Data_Type >::NCells = 0u
```

Definition at line 28 of file barray-bones.hpp.

7.1.5.9 visited

```
template<typename Cell_Type = bool, typename Data_Type = bool>
bool BArray< Cell_Type, Data_Type >::visited = false
```

This is as a reference, if we need to iterate through the cells and we need to keep track which were visited, we use this as a reference. So that if cell.visited = true and visited = true, it means that we haven't been here yet. Ideally, any routine using this->visited should switch it at the beginning of the routine.

Definition at line 43 of file barray-bones.hpp.

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

include/barry/barray-bones.hpp

7.2 BArrayCell< Cell_Type, Data_Type > Class Template Reference

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

Public Member Functions

- BArrayCell (BArray < Cell Type, Data Type > *Array , uint i , uint j , bool check bounds=true)
- ∼BArrayCell ()
- void operator= (const Cell_Type &val)
- void operator+= (const Cell_Type &val)
- void operator-= (const Cell_Type &val)
- void operator*= (const Cell_Type &val)
- void operator/= (const Cell_Type &val)
- operator Cell_Type () const
- bool operator== (const Cell_Type &val) const

7.2.1 Detailed Description

```
template<typename Cell_Type = bool, typename Data_Type = bool> class BArrayCell< Cell_Type, Data_Type >
```

Definition at line 7 of file barraycell-bones.hpp.

7.2.2 Constructor & Destructor Documentation

7.2.2.1 BArrayCell()

Definition at line 16 of file barraycell-bones.hpp.

7.2.2.2 ~BArrayCell()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
BArrayCell< Cell_Type, Data_Type >::~BArrayCell ( ) [inline]
```

Definition at line 28 of file barraycell-bones.hpp.

7.2.3 Member Function Documentation

7.2.3.1 operator Cell_Type()

```
template<typename Cell_Type , typename Data_Type >
BArrayCell< Cell_Type, Data_Type >::operator Cell_Type [inline]
```

Definition at line 58 of file barraycell-meat.hpp.

7.2.3.2 operator*=()

Definition at line 40 of file barraycell-meat.hpp.

7.2.3.3 operator+=()

Definition at line 18 of file barraycell-meat.hpp.

7.2.3.4 operator-=()

Definition at line 29 of file barraycell-meat.hpp.

7.2.3.5 operator/=()

Definition at line 49 of file barraycell-meat.hpp.

7.2.3.6 operator=()

Definition at line 7 of file barraycell-meat.hpp.

7.2.3.7 operator==()

Definition at line 63 of file barraycell-meat.hpp.

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

- include/barry/barraycell-bones.hpp
- include/barry/barraycell-meat.hpp

7.3 BArrayCell_const< Cell_Type, Data_Type > Class Template Reference

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

Public Member Functions

- BArrayCell_const (const BArray< Cell_Type, Data_Type > *Array_, uint i_, uint i_, bool check_bounds=true)
- ∼BArrayCell const ()
- operator Cell_Type () const
- bool operator== (const Cell_Type &val) const
- bool operator!= (const Cell_Type &val) const
- bool operator< (const Cell_Type &val) const
- bool operator> (const Cell_Type &val) const
- bool operator<= (const Cell_Type &val) const
- bool operator>= (const Cell_Type &val) const

7.3.1 Detailed Description

```
template<typename Cell_Type = bool, typename Data_Type = bool> class BArrayCell_const< Cell_Type, Data_Type >
```

Definition at line 41 of file barraycell-bones.hpp.

7.3.2 Constructor & Destructor Documentation

7.3.2.1 BArrayCell_const()

Definition at line 50 of file barraycell-bones.hpp.

7.3.2.2 ~BArrayCell const()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
BArrayCell_const < Cell_Type, Data_Type >::~BArrayCell_const ( ) [inline]
```

Definition at line 62 of file barraycell-bones.hpp.

7.3.3 Member Function Documentation

7.3.3.1 operator Cell_Type()

```
template<typename Cell_Type , typename Data_Type >
BArrayCell_const< Cell_Type, Data_Type >::operator Cell_Type [inline]
```

Definition at line 68 of file barraycell-meat.hpp.

7.3.3.2 operator"!=()

Definition at line 78 of file barraycell-meat.hpp.

7.3.3.3 operator<()

Definition at line 83 of file barraycell-meat.hpp.

7.3.3.4 operator<=()

Definition at line 93 of file barraycell-meat.hpp.

7.3.3.5 operator==()

Definition at line 73 of file barraycell-meat.hpp.

7.3.3.6 operator>()

Definition at line 88 of file barraycell-meat.hpp.

7.3.3.7 operator>=()

Definition at line 98 of file barraycell-meat.hpp.

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

- include/barry/barraycell-bones.hpp
- include/barry/barraycell-meat.hpp

7.4 Cell< Cell_Type > Class Template Reference

Entries in BArray. For now, it only has two members:

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

Public Member Functions

```
• Cell ()
```

- Cell (Cell_Type value_, bool visited_=false)
- ~Cell ()
- Cell (const Cell< Cell_Type > &arg)
- Cell< Cell_Type > & operator= (Cell< Cell_Type > &other)
- Cell (Cell< Cell_Type > &&arg) noexcept
- Cell< Cell_Type > & operator= (Cell< Cell_Type > &&other) noexcept
- void add (Cell Type x)
- operator Cell_Type () const
- void add (double x)
- void add (unsigned int x)
- void add (int x)
- Cell ()
- Cell ()
- Cell ()

Public Attributes

- Cell_Type value
- bool visited

7.4.1 Detailed Description

```
template < class Cell_Type > class Cell < Cell_Type >
```

Entries in BArray. For now, it only has two members:

- · value: the content
- · visited: boolean (just a convenient)

Definition at line 13 of file cell-bones.hpp.

7.4.2 Constructor & Destructor Documentation

7.4.2.1 Cell() [1/7]

```
template<class Cell_Type >
Cell< Cell_Type >::Cell ( )
```

7.4.2.2 Cell() [2/7]

Definition at line 18 of file cell-bones.hpp.

7.4.2.3 ∼CeII()

```
template<class Cell_Type >
Cell< Cell_Type >::~Cell ( ) [inline]
```

Definition at line 20 of file cell-bones.hpp.

7.4.2.4 Cell() [3/7]

Definition at line 24 of file cell-bones.hpp.

7.4.2.5 Cell() [4/7]

Definition at line 30 of file cell-bones.hpp.

7.4.2.6 Cell() [5/7]

```
Cell< double >::Cell ( ) [inline]
```

Definition at line 44 of file cell-meat.hpp.

7.4.2.7 Cell() [6/7]

```
Cell< uint >::Cell ( ) [inline]
```

Definition at line 45 of file cell-meat.hpp.

7.4.2.8 Cell() [7/7]

```
Cell< int >::Cell ( ) [inline]
```

Definition at line 46 of file cell-meat.hpp.

7.4.3 Member Function Documentation

7.4.3.1 add() [1/4]

7.4.3.2 add() [2/4]

Definition at line 24 of file cell-meat.hpp.

7.4.3.3 add() [3/4]

```
void Cell< int >::add (
          int x ) [inline]
```

Definition at line 34 of file cell-meat.hpp.

7.4.3.4 add() [4/4]

```
void Cell< unsigned int >:: add ( unsigned int x ) [inline]
```

Definition at line 29 of file cell-meat.hpp.

7.4.3.5 operator Cell_Type()

```
template<class Cell_Type >
Cell< Cell_Type >::operator Cell_Type ( ) const [inline]
```

Definition at line 41 of file cell-bones.hpp.

7.4.3.6 operator=() [1/2]

Definition at line 14 of file cell-meat.hpp.

7.4.3.7 operator=() [2/2]

Definition at line 7 of file cell-meat.hpp.

7.4.4 Member Data Documentation

7.4.4.1 value

```
template<class Cell_Type >
Cell_Type Cell< Cell_Type >::value
```

Definition at line 15 of file cell-bones.hpp.

7.4.4.2 visited

```
template<class Cell_Type >
bool Cell< Cell_Type >::visited
```

Definition at line 16 of file cell-bones.hpp.

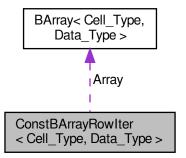
The documentation for this class was generated from the following files:

- include/barry/cell-bones.hpp
- include/barry/cell-meat.hpp

7.5 ConstBArrayRowlter< Cell_Type, Data_Type > Class Template Reference

```
#include <barray-iterator.hpp>
```

Collaboration diagram for ConstBArrayRowIter< Cell_Type, Data_Type >:



Public Member Functions

- ConstBArrayRowlter (const BArray< Cell_Type, Data_Type > *Array_)
- \sim ConstBArrayRowIter ()

Public Attributes

- uint current_row
- uint current_col
- Row_type< Cell_Type >::const_iterator iter
- const BArray< Cell_Type, Data_Type > * Array

7.5.1 Detailed Description

```
template<typename Cell_Type, typename Data_Type> class ConstBArrayRowlter< Cell_Type, Data_Type>
```

Definition at line 10 of file barray-iterator.hpp.

7.5.2 Constructor & Destructor Documentation

7.5.2.1 ConstBArrayRowIter()

Definition at line 17 of file barray-iterator.hpp.

7.5.2.2 ∼ConstBArrayRowlter()

```
template<typename Cell_Type , typename Data_Type >
ConstBArrayRowIter< Cell_Type, Data_Type >::~ConstBArrayRowIter ( ) [inline]
```

Definition at line 29 of file barray-iterator.hpp.

7.5.3 Member Data Documentation

7.5.3.1 Array

```
template<typename Cell_Type , typename Data_Type >
const BArray<Cell_Type,Data_Type>* ConstBArrayRowIter< Cell_Type, Data_Type >::Array
```

Definition at line 15 of file barray-iterator.hpp.

7.5.3.2 current_col

```
template<typename Cell_Type , typename Data_Type >
uint ConstBArrayRowIter< Cell_Type, Data_Type >::current_col
```

Definition at line 13 of file barray-iterator.hpp.

7.5.3.3 current_row

```
template<typename Cell_Type , typename Data_Type >
uint ConstBArrayRowIter< Cell_Type, Data_Type >::current_row
```

Definition at line 13 of file barray-iterator.hpp.

7.5.3.4 iter

```
template<typename Cell_Type , typename Data_Type >
Row_type<Cell_Type>::const_iterator ConstBArrayRowIter< Cell_Type, Data_Type >::iter
```

Definition at line 14 of file barray-iterator.hpp.

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

• include/barry/barray-iterator.hpp

7.6 Counter< Array_Type, Data_Type > Class Template Reference

A counter function based on change statistics.

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

Public Member Functions

- Counter< Array_Type, Data_Type > operator= (const Counter< Array_Type, Data_Type > &counter_)
- ∼Counter ()
- double count (Array_Type &Array, uint i, uint j)
- double init (Array_Type &Array, uint i, uint j)

Creator passing a counter and an initializer

Parameters

count_fun←	The main counter function.
init_fun_	The initializer function can also be used to check if the BArray as the needed variables (see BArray::data).
data_	Data to be used with the counter.
delete_← data_	When true, the destructor will delete the pointer in the main data.

- Counter ()
- Counter (Counter_fun_type
 Array_Type, Data_Type > count_fun_, Counter_fun_type
 Array_Type, Data_Type > init_fun_=nullptr, Data_Type *data_=nullptr, bool delete_data_=false)

Counter (const Counter< Array_Type, Data_Type > &counter_)

Public Attributes

```
    Counter_fun_type< Array_Type, Data_Type > count_fun
    Counter_fun_type< Array_Type, Data_Type > init_fun
```

Data_Type * data = nullptrbool delete_data = false

7.6.1 Detailed Description

```
template<typename Array_Type = BArray<>, typename Data_Type = bool> class Counter< Array_Type, Data_Type >
```

A counter function based on change statistics.

This class is used by CountStats and StatsCounter as a way to count statistics using change statistics.

Definition at line 38 of file counters-bones.hpp.

7.6.2 Constructor & Destructor Documentation

7.6.2.1 Counter() [1/3]

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Counter< Array_Type, Data_Type >::Counter ( ) [inline]
```

Definition at line 57 of file counters-bones.hpp.

7.6.2.2 Counter() [2/3]

Definition at line 59 of file counters-bones.hpp.

7.6.2.3 Counter() [3/3]

Definition at line 7 of file counters-meat.hpp.

7.6.2.4 ~Counter()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Counter< Array_Type, Data_Type >::~Counter ( ) [inline]
```

Definition at line 71 of file counters-bones.hpp.

7.6.3 Member Function Documentation

7.6.3.1 count()

Definition at line 100 of file counters-meat.hpp.

7.6.3.2 init()

Definition at line 108 of file counters-meat.hpp.

7.6.3.3 operator=()

Definition at line 24 of file counters-meat.hpp.

7.6.4 Member Data Documentation

7.6.4.1 count_fun

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Counter_fun_type<Array_Type,Data_Type> Counter< Array_Type, Data_Type>::count_fun
```

Definition at line 41 of file counters-bones.hpp.

7.6.4.2 data

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Data_Type* Counter< Array_Type, Data_Type >::data = nullptr
```

Definition at line 43 of file counters-bones.hpp.

7.6.4.3 delete_data

```
template<typename Array_Type = BArray<>>, typename Data_Type = bool>
bool Counter< Array_Type, Data_Type >::delete_data = false
```

Definition at line 44 of file counters-bones.hpp.

7.6.4.4 init fun

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Counter_fun_type<Array_Type, Data_Type> Counter< Array_Type, Data_Type>:::nit_fun
```

Definition at line 42 of file counters-bones.hpp.

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

- include/barry/counters-bones.hpp
- include/barry/counters-meat.hpp

7.7 Counters < Array_Type, Data_Type > Class Template Reference

Vector of counters.

#include <counters-bones.hpp>

Public Member Functions

- · Counters ()
- ∼Counters ()
- Counters (const Counters < Array_Type, Data_Type > &counter_)
- Counters< Array_Type, Data_Type > operator= (const Counters< Array_Type, Data_Type > &counter_)
- Counter< Array_Type, Data_Type > & operator[] (uint idx)

Returns a pointer to a particular counter.

• std::size t size () const noexcept

Number of counters in the set.

- void add_counter (Counter< Array_Type, Data_Type > &counter)
- void add_counter (Counter< Array_Type, Data_Type > *counter)
- void add_counter (Counter_fun_type< Array_Type, Data_Type > count_fun_, Counter_fun_type< Array_
 —
 Type, Data_Type > init_fun_=nullptr, Data_Type *data_=nullptr, bool delete_data_=false)
- void clear ()

7.7.1 Detailed Description

```
\label{template} $$ $ template < typename \ Array_Type = BArray <>, typename \ Data_Type = bool > class \ Counters < Array_Type, \ Data_Type > $$
```

Vector of counters.

Various functions hold more than one counter, so this class is a helper class that allows managing multiple counters efficiently. The main data is a vector to pointers of counters.

Definition at line 92 of file counters-bones.hpp.

7.7.2 Constructor & Destructor Documentation

7.7.2.1 Counters() [1/2]

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Counters< Array_Type, Data_Type >::Counters ( ) [inline]
```

Definition at line 101 of file counters-bones.hpp.

7.7.2.2 ~Counters()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Counters< Array_Type, Data_Type >::~Counters () [inline]
```

Definition at line 104 of file counters-bones.hpp.

7.7.2.3 Counters() [2/2]

Definition at line 48 of file counters-meat.hpp.

7.7.3 Member Function Documentation

7.7.3.1 add_counter() [1/3]

Definition at line 121 of file counters-meat.hpp.

7.7.3.2 add_counter() [2/3]

Definition at line 132 of file counters-meat.hpp.

7.7.3.3 add_counter() [3/3]

Definition at line 142 of file counters-meat.hpp.

7.7.3.4 clear()

```
template<typename Array_Type , typename Data_Type >
void Counters< Array_Type, Data_Type >::clear [inline]
```

Definition at line 166 of file counters-meat.hpp.

7.7.3.5 operator=()

Definition at line 72 of file counters-meat.hpp.

7.7.3.6 operator[]()

Returns a pointer to a particular counter.

Parameters

```
idx Id of the counter
```

Returns

```
Counter<Array_Type,Data_Type>*
```

Definition at line 116 of file counters-meat.hpp.

7.7.3.7 size()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
std::size_t Counters< Array_Type, Data_Type >::size ( ) const [inline], [noexcept]
```

Number of counters in the set.

Returns

uint

Definition at line 124 of file counters-bones.hpp.

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

- include/barry/counters-bones.hpp
- include/barry/counters-meat.hpp

7.8 Entries < Cell_Type > Class Template Reference

A wrapper class to store source, target, val from a BArray object.

```
#include <typedefs.hpp>
```

Public Member Functions

- Entries ()
- Entries (uint n)
- ∼Entries ()
- void resize (uint n)

Public Attributes

- std::vector< uint > source
- std::vector< uint > target
- std::vector< Cell_Type > val

7.8.1 Detailed Description

```
\label{template} \begin{tabular}{ll} template < typename Cell_Type > \\ class Entries < Cell_Type > \\ \end{tabular}
```

A wrapper class to store source, target, val from a BArray object.

Template Parameters

```
Cell_Type Any type
```

Definition at line 70 of file typedefs.hpp.

7.8.2 Constructor & Destructor Documentation

7.8.2.1 Entries() [1/2]

```
template<typename Cell_Type >
Entries< Cell_Type >::Entries ( ) [inline]
```

Definition at line 76 of file typedefs.hpp.

7.8.2.2 Entries() [2/2]

```
template<typename Cell_Type >
Entries< Cell_Type >::Entries (
          uint n) [inline]
```

Definition at line 77 of file typedefs.hpp.

7.8.2.3 \sim Entries()

```
template<typename Cell_Type >
Entries< Cell_Type >::~Entries ( ) [inline]
```

Definition at line 84 of file typedefs.hpp.

7.8.3 Member Function Documentation

7.8.3.1 resize()

Definition at line 86 of file typedefs.hpp.

7.8.4 Member Data Documentation

7.8.4.1 source

```
template<typename Cell_Type >
std::vector< uint > Entries< Cell_Type >::source
```

Definition at line 72 of file typedefs.hpp.

7.8.4.2 target

```
template<typename Cell_Type >
std::vector< uint > Entries< Cell_Type >::target
```

Definition at line 73 of file typedefs.hpp.

7.9 Flock Class Reference 61

7.8.4.3 val

```
template<typename Cell_Type >
std::vector< Cell_Type > Entries< Cell_Type >::val
```

Definition at line 74 of file typedefs.hpp.

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

• include/barry/typedefs.hpp

7.9 Flock Class Reference

A Flock is a group of Geese.

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

Public Member Functions

- Flock ()
- ∼Flock ()
- unsigned int add_data (std::vector< std::vector< unsigned int > > &annotations, std::vector< unsigned int > &geneid, std::vector< int > &parent, std::vector< bool > &duplication)

Add a tree to the flock.

• void set seed (const unsigned int &s)

Set the seed of the model.

- void init ()
- phylocounters::PhyloCounters * counters_ptr ()

Returns the joint likelihood of the model.

Information about the model

- · unsigned int nfuns () const noexcept
- unsigned int ntrees () const noexcept
- std::vector< unsigned int > nnodes () const noexcept
- std::vector< unsigned int > nleafs () const noexcept
- unsigned int nterms () const

Public Attributes

- std::vector< Geese > dat
- unsigned int nfunctions = 0u
- bool initialized = false
- std::mt19937 rengine
- phylocounters::PhyloModel support = phylocounters::PhyloModel()

7.9.1 Detailed Description

A Flock is a group of Geese.

This object buils a model with multiple trees (Geese objects), with all of these using the same PhyloModel object. Available counters (terms) can be found in counter-phylo.

Definition at line 14 of file flock-bones.hpp.

7.9.2 Constructor & Destructor Documentation

7.9.2.1 Flock()

```
Flock::Flock ( ) [inline]
```

Definition at line 25 of file flock-bones.hpp.

7.9.2.2 ∼Flock()

```
Flock::~Flock ( ) [inline]
```

Definition at line 26 of file flock-bones.hpp.

7.9.3 Member Function Documentation

7.9.3.1 add_data()

```
unsigned int Flock::add_data (
    std::vector< std::vector< unsigned int > & annotations,
    std::vector< unsigned int > & geneid,
    std::vector< int > & parent,
    std::vector< bool > & duplication ) [inline]
```

Add a tree to the flock.

Parameters

annotations	see Geese::Geese.
geneid	see Geese.
parent	see Geese.
duplication	see Geese.

7.9 Flock Class Reference 63

Returns

unsigned int The number of tree in the model (starting from zero).

Definition at line 6 of file flock-meet.hpp.

7.9.3.2 counters_ptr()

```
phylocounters::PhyloCounters * Flock::counters_ptr ( ) [inline]
```

Definition at line 75 of file flock-meet.hpp.

7.9.3.3 init()

```
void Flock::init ( ) [inline]
```

Definition at line 41 of file flock-meet.hpp.

7.9.3.4 likelihood_joint()

Returns the joint likelihood of the model.

Parameters

par	Vector of model parameters.	
as_log	When true it will return the value as log.	
use_likelihood_sequence When true (default) will compute the likelihood using the reduced sequence which is faster.		

Returns

double

Definition at line 84 of file flock-meet.hpp.

7.9.3.5 nfuns()

```
unsigned int Flock::nfuns ( ) const [inline], [noexcept]
```

Definition at line 109 of file flock-meet.hpp.

7.9.3.6 nleafs()

```
std::vector< unsigned int > Flock::nleafs ( ) const [inline], [noexcept]
```

Definition at line 132 of file flock-meet.hpp.

7.9.3.7 nnodes()

```
std::vector< unsigned int > Flock::nnodes ( ) const [inline], [noexcept]
```

Definition at line 121 of file flock-meet.hpp.

7.9.3.8 nterms()

```
unsigned int Flock::nterms ( ) const [inline]
```

Definition at line 144 of file flock-meet.hpp.

7.9.3.9 ntrees()

```
unsigned int Flock::ntrees ( ) const [inline], [noexcept]
```

Definition at line 115 of file flock-meet.hpp.

7.9.3.10 set_seed()

Set the seed of the model.

7.9 Flock Class Reference 65

Parameters

s Passed to the rengine.seed() member object.

Definition at line 37 of file flock-meet.hpp.

7.9.4 Member Data Documentation

7.9.4.1 dat

std::vector< Geese > Flock::dat

Definition at line 17 of file flock-bones.hpp.

7.9.4.2 initialized

bool Flock::initialized = false

Definition at line 19 of file flock-bones.hpp.

7.9.4.3 nfunctions

unsigned int Flock::nfunctions = 0u

Definition at line 18 of file flock-bones.hpp.

7.9.4.4 rengine

std::mt19937 Flock::rengine

Definition at line 22 of file flock-bones.hpp.

7.9.4.5 support

```
phylocounters::PhyloModel Flock::support = phylocounters::PhyloModel()
```

Definition at line 23 of file flock-bones.hpp.

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

- include/barry/models/geese/flock-bones.hpp
- include/barry/models/geese/flock-meet.hpp

7.10 FreqTable < T > Class Template Reference

Database of statistics.

```
#include <statsdb.hpp>
```

Public Member Functions

- FreqTable ()
- ∼FreqTable ()
- void add (const std::vector< T > &x)
- Counts_type as_vector () const
- MapVec_type< T, uint > get_data () const
- const MapVec_type< T, uint > * get_data_ptr () const
- void clear ()
- void reserve (unsigned int n)
- void print () const
- size_t size () const noexcept

7.10.1 Detailed Description

```
template<typename T = double> class FreqTable< T >
```

Database of statistics.

This is mostly used in Support.

Definition at line 16 of file statsdb.hpp.

7.10.2 Constructor & Destructor Documentation

7.10.2.1 FreqTable()

```
template<typename T = double>
FreqTable< T >::FreqTable ( ) [inline]
```

Definition at line 28 of file statsdb.hpp.

7.10.2.2 \sim FreqTable()

```
template<typename T = double>
FreqTable< T >::~FreqTable ( ) [inline]
```

Definition at line 29 of file statsdb.hpp.

7.10.3 Member Function Documentation

7.10.3.1 add()

Definition at line 47 of file statsdb.hpp.

7.10.3.2 as_vector()

```
template<typename T >
Counts_type FreqTable< T >::as_vector [inline]
```

Definition at line 61 of file statsdb.hpp.

7.10.3.3 clear()

```
template<typename T >
void FreqTable< T >::clear [inline]
```

Definition at line 83 of file statsdb.hpp.

7.10.3.4 get_data()

```
template<typename T >
MapVec_type< T, uint > FreqTable< T >::get_data [inline]
```

Definition at line 73 of file statsdb.hpp.

7.10.3.5 get_data_ptr()

```
template<typename T >
const MapVec_type< T, uint > * FreqTable< T >::get_data_ptr [inline]
```

Definition at line 78 of file statsdb.hpp.

7.10.3.6 print()

```
template<typename T >
void FreqTable< T >::print [inline]
```

Definition at line 102 of file statsdb.hpp.

7.10.3.7 reserve()

```
\label{template} \begin{tabular}{ll} template < typename T > \\ void FreqTable < T > :: reserve ( \\ & unsigned int $n$ ) [inline] \end{tabular}
```

Definition at line 89 of file statsdb.hpp.

7.10.3.8 size()

```
template<typename T >
size_t FreqTable< T >::size [inline], [noexcept]
```

Definition at line 113 of file statsdb.hpp.

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

• include/barry/statsdb.hpp

7.11 Geese Class Reference

Annotated Phylo Model.

#include <geese-bones.hpp>

Public Member Functions

- ∼Geese ()
- void init ()
- void inherit_support (const Geese &model_, bool delete_support_=false)
- void calc sequence (Node *n=nullptr)
- void calc_likelihood_sequence ()
- double likelihood (const std::vector< double > &par, bool as log=false, bool use likelihood sequence=true)
- double likelihood_exhaust (const std::vector< double > &par)
- std::vector< double > get_probabilities () const
- · void set seed (const unsigned int &s)
- std::vector< std::vector< unsigned int > > simulate (const std::vector< double > &par)
- std::vector< std::vector< double >> observed counts ()
- void print observed counts ()
- std::vector< std::vector< double > > predict (const std::vector< double > &p, std::vector< std::vector< double > > *res_prob=nullptr)

Calculate the conditional probability.

- void init node (Node &n)
- void update annotations (unsigned int nodeid, std::vector< unsigned int > newann)

Construct a new Geese object

The model includes a total of N + 1 nodes, the + 1 beign the root node.

Parameters

annotations	A vector of vectors with annotations. It should be of length k (number of functions). Each vector should be of length $\mathbb N$ (equal to the number of nodes, including interior). Possible values are 0, 1, and 9.
geneid	Id of the gene. It should be of length ${\tt N}.$
parent	Id of the parent gene. Also of length N

- Geese ()
- Geese (std::vector< std::vector< unsigned int > > &annotations, std::vector< unsigned int > &geneid, std::vector< int > &parent, std::vector< bool > &duplication)
- Geese (const Geese &model_, bool copy_data=true)
- Geese (Geese &&x) noexcept
- Geese & operator= (const Geese &model_)=delete
- Geese & operator= (Geese &&model_) noexcept=delete

Information about the model

- · unsigned int nfuns () const noexcept
- unsigned int nnodes () const noexcept
- unsigned int nleafs () const noexcept
- · unsigned int nterms () const

Public Attributes

- · unsigned int nfunctions
- barry::Map< unsigned int, Node > nodes
- barry::MapVec_type< unsigned int > map_to_nodes
- std::vector< unsigned int > sequence
- std::vector< unsigned int > likelihood sequence
- bool initialized = false
- bool delete rengine = false
- bool delete_counters = false
- bool delete_support = false

Shared objects within a <tt>Geese</tt>

Since users may start adding counters before initializing the PhyloModel object, the object counter is initialized first.

While the member support has an rengine, since Geese can sample trees, we have the option to keep it separate.

```
• std::mt19937 * rengine = nullptr
```

- phylocounters::PhyloCounters * counters = nullptr
- phylocounters::PhyloModel * support = nullptr
- std::vector< std::vector< bool >> states

7.11.1 Detailed Description

Annotated Phylo Model.

A list of available terms for this model can be found in the Phylo counters section.

Definition at line 70 of file geese-bones.hpp.

7.11.2 Constructor & Destructor Documentation

7.11.2.1 Geese() [1/4]

```
Geese::Geese ( ) [inline]
```

Definition at line 6 of file geese-meat-constructors.hpp.

7.11.2.2 Geese() [2/4]

Definition at line 17 of file geese-meat-constructors.hpp.

7.11.2.3 Geese() [3/4]

Definition at line 157 of file geese-meat-constructors.hpp.

7.11.2.4 Geese() [4/4]

Definition at line 232 of file geese-meat-constructors.hpp.

7.11.2.5 ∼Geese()

```
Geese::~Geese ( ) [inline]
```

Definition at line 71 of file geese-meat.hpp.

7.11.3 Member Function Documentation

7.11.3.1 calc_likelihood_sequence()

```
void Geese::calc_likelihood_sequence ( ) [inline]
```

Definition at line 234 of file geese-meat.hpp.

7.11.3.2 calc_sequence()

```
void Geese::calc_sequence (
          Node * n = nullptr ) [inline]
```

Definition at line 191 of file geese-meat.hpp.

7.11.3.3 get_probabilities()

```
std::vector< double > Geese::get_probabilities ( ) const [inline]
```

Definition at line 271 of file geese-meat.hpp.

7.11.3.4 inherit_support()

Definition at line 140 of file geese-meat.hpp.

7.11.3.5 init()

```
void Geese::init ( ) [inline]
```

Definition at line 83 of file geese-meat.hpp.

7.11.3.6 init_node()

```
void Geese::init_node ( \begin{tabular}{ll} Node & n \end{tabular} \begin{tabular}{ll} [inline] \end{tabular}
```

Definition at line 6 of file geese-meat.hpp.

7.11.3.7 likelihood()

Definition at line 6 of file geese-meat-likelihood.hpp.

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7.11.3.8 likelihood_exhaust()

Definition at line 7 of file geese-meat-likelihood_exhaust.hpp.

7.11.3.9 nfuns()

```
unsigned int Geese::nfuns ( ) const [inline], [noexcept]
```

Definition at line 287 of file geese-meat.hpp.

7.11.3.10 nleafs()

```
unsigned int Geese::nleafs ( ) const [inline], [noexcept]
```

Definition at line 295 of file geese-meat.hpp.

7.11.3.11 nnodes()

```
unsigned int Geese::nnodes ( ) const [inline], [noexcept]
```

Definition at line 291 of file geese-meat.hpp.

7.11.3.12 nterms()

```
unsigned int Geese::nterms ( ) const [inline]
```

Definition at line 305 of file geese-meat.hpp.

7.11.3.13 observed_counts()

```
\verb|std::vector| < \verb|std::vector| < \verb|double| > > Geese::observed_counts () [inline]|
```

Definition at line 312 of file geese-meat.hpp.

7.11.3.14 operator=() [1/2]

7.11.3.15 operator=() [2/2]

```
Geese& Geese::operator= (
          Geese && model_ ) [delete], [noexcept]
```

7.11.3.16 predict()

Calculate the conditional probability.

Parameters

```
p Vector of parameters
```

Returns

```
std::vector< double > Returns the posterior probability
```

Definition at line 7 of file geese-meat-predict.hpp.

7.11.3.17 print_observed_counts()

```
void Geese::print_observed_counts ( ) [inline]
```

Definition at line 360 of file geese-meat.hpp.

7.11.3.18 set_seed()

Definition at line 8 of file geese-meat-simulate.hpp.

7.11 Geese Class Reference 75

7.11.3.19 simulate()

```
std::vector< std::vector< unsigned int >> Geese::simulate ( const std::vector< double > & par ) [inline]
```

Definition at line 12 of file geese-meat-simulate.hpp.

7.11.3.20 update_annotations()

```
void Geese::update_annotations (
          unsigned int nodeid,
          std::vector< unsigned int > newann ) [inline]
```

Definition at line 168 of file geese-meat.hpp.

7.11.4 Member Data Documentation

7.11.4.1 counters

```
phylocounters::PhyloCounters* Geese::counters = nullptr
```

Definition at line 85 of file geese-bones.hpp.

7.11.4.2 delete_counters

```
bool Geese::delete_counters = false
```

Definition at line 102 of file geese-bones.hpp.

7.11.4.3 delete_rengine

```
bool Geese::delete_rengine = false
```

Definition at line 101 of file geese-bones.hpp.

7.11.4.4 delete_support

```
bool Geese::delete_support = false
```

Definition at line 103 of file geese-bones.hpp.

7.11.4.5 initialized

```
bool Geese::initialized = false
```

Definition at line 100 of file geese-bones.hpp.

7.11.4.6 likelihood_sequence

```
std::vector< unsigned int > Geese::likelihood_sequence
```

Definition at line 97 of file geese-bones.hpp.

7.11.4.7 map_to_nodes

```
barry::MapVec_type< unsigned int > Geese::map_to_nodes
```

Definition at line 93 of file geese-bones.hpp.

7.11.4.8 nfunctions

```
unsigned int Geese::nfunctions
```

Definition at line 91 of file geese-bones.hpp.

7.11.4.9 nodes

```
barry::Map< unsigned int, Node > Geese::nodes
```

Definition at line 92 of file geese-bones.hpp.

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7.11.4.10 rengine

```
std::mt19937* Geese::rengine = nullptr
```

Definition at line 84 of file geese-bones.hpp.

7.11.4.11 sequence

```
std::vector< unsigned int > Geese::sequence
```

Definition at line 96 of file geese-bones.hpp.

7.11.4.12 states

```
std::vector< std::vector< bool > > Geese::states
```

Definition at line 87 of file geese-bones.hpp.

7.11.4.13 support

```
phylocounters::PhyloModel* Geese::support = nullptr
```

Definition at line 86 of file geese-bones.hpp.

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

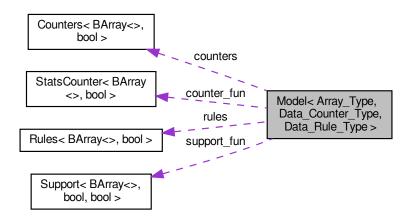
- include/barry/models/geese/geese-bones.hpp
- include/barry/models/geese/geese-meat-constructors.hpp
- include/barry/models/geese/geese-meat-likelihood.hpp
- include/barry/models/geese/geese-meat-likelihood_exhaust.hpp
- include/barry/models/geese/geese-meat-predict.hpp
- include/barry/models/geese/geese-meat-simulate.hpp
- include/barry/models/geese/geese-meat.hpp

7.12 Model < Array_Type, Data_Counter_Type, Data_Rule_Type > Class Template Reference

General framework for discrete exponential models. This class allows generating discrete exponential models in the form of a linear exponential model:

#include <model-bones.hpp>

Collaboration diagram for Model < Array_Type, Data_Counter_Type, Data_Rule_Type >:



Public Member Functions

- Model ()
- Model (uint size)
- Model (const Model < Array_Type, Data_Counter_Type, Data_Rule_Type > &Model_)
- Model Array_Type, Data_Counter_Type, Data_Rule_Type > & operator= (const Model Array_Type, Data_Counter_Type, Data_Rule_Type > &Model_)
- ∼Model ()
- void store_psets () noexcept
- void set keygen (std::function < std::vector < double > (const Array Type &) > keygen)
- uint add_array (const Array_Type &Array_, bool force_new=false)

Adds an array to the support of not already included.

- void print_stats (uint i) const
- Array_Type sample (const Array_Type &Array_, const std::vector< double > ¶ms={})
- Array_Type sample (const uint &i, const std::vector< double > ¶ms)

Wrappers for the <tt>Counters</tt> member.

These will add counters to the model, which are shared by the support and the actual counter function.

- void add_counter (Counter< Array_Type, Data_Counter_Type > &counter)
- void add_counter (Counter< Array_Type, Data_Counter_Type > *counter)
- void add_counter (Counter_fun_type < Array_Type, Data_Counter_Type > count_fun_, Counter_fun_type <
 Array_Type, Data_Counter_Type > init_fun_=nullptr, Data_Counter_Type *data_=nullptr, bool delete_
 data_=false)

void set_counters (Counters < Array_Type, Data_Counter_Type > *counters_)

Wrappers for the <tt>Rules</tt> member.

These will add rules to the model, which are shared by the support and the actual counter function.

- void add_rule (Rule < Array_Type, Data_Rule_Type > &rule)
- void add rule (Rule < Array Type, Data Rule Type > *rule)
- void add_rule (Rule_fun_type< Array_Type, Data_Rule_Type > count_fun_, Data_Rule_Type *data_←
 =nullptr, bool delete data =false)
- void set rules (Rules < Array Type, Data Rule Type > *rules)

Likelihood functions.

Calculation of likelihood functions is done reusing normalizing constants. Before recalculating the normalizing constant, the function checks whether params matches the last set vector of parameters used to compute it.

Parameters

params	Vector of parameters	
as_log	When true, the function returns the log-likelihood.	

- double likelihood (const std::vector< double > ¶ms, const uint &i, bool as log=false)
- double likelihood (const std::vector< double > ¶ms, const std::vector< double > &target_, const uint &i, bool as log=false)
- double likelihood total (const std::vector< double > ¶ms, bool as log=false)

Extract elements by index

Parameters

i	Index relative to the array in the model.	
params	A new vector of model parameters to compute the normalizing constant.	
as_log	When true returns the logged version of the normalizing constant.	

- double get_norm_const (const std::vector< double > ¶ms, const uint &i, bool as_log=false)
- const std::vector< Array_Type > * get_pset (const uint &i)
- const std::vector< std::vector< double > > * get_stats (const uint &i)

Size of the model

Number of different supports included in the model

This will return the size of stats.

Returns

size() returns the number of arrays in the model.
size_unique() returns the number of unique arrays (according to the hasher) in the model.
nterms() returns the number of terms in the model.

- unsigned int size () const noexcept
- unsigned int size_unique () const noexcept
- unsigned int nterms () const noexcept

Public Attributes

- std::vector < Counts_type > stats
- std::vector< uint > n_arrays_per_stats
- MapVec_type< double, uint > keys2support

Map of types of arrays to support sets.

std::vector< std::vector< double >> params last

Vector of the previously used parameters.

- std::vector< double > normalizing constants
- std::vector< bool > first_calc_done
- std::function< std::vector< double >const Array Type &)> keygen = nullptr

Function to extract features of the array to be hash.

Container space for the powerset (and its sufficient stats)

This is useful in the case of using simulations or evaluating functions that need to account for the full set of states.

- bool with pset = false
- std::vector< std::vector< Array_Type >> pset_arrays
- std::vector< std::vector< double >>> pset_stats
- std::vector< std::vector< double >> pset_probs

Information about the arrays used in the model

target_stats holds the observed sufficient statistics for each array in the dataset. array_frequency contains the frequency with which each of the target stats (arrays) shows in the support. array2support maps array indices (0, 1, ...) to the corresponding support.

- std::vector< std::vector< double > > target stats
- std::vector< uint > array_frequency
- std::vector< uint > arrays2support

Functions to compute statistics

Arguments are recycled to save memory and computation.

- Counters
 Array_Type, Data_Counter_Type > counters
- Rules < Array_Type, Data_Rule_Type > rules
- Support < Array Type, Data Counter Type, Data Rule Type > support fun
- StatsCounter< Array_Type, Data_Counter_Type > counter_fun

Random number generation

Random number generation

- std::mt19937 * rengine = nullptr
- bool delete_rengine = false
- void set_rengine (std::mt19937 *rengine_, bool delete_=false)
- void set_seed (unsigned int s)

7.12.1 Detailed Description

template<typename Array_Type = BArray<>>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool> class Model< Array_Type, Data_Counter_Type, Data_Rule_Type >

General framework for discrete exponential models. This class allows generating discrete exponential models in the form of a linear exponential model:

$$\frac{\exp\left(\theta^{\dagger}c(A)\right)}{\sum_{A'\in\mathcal{A}}\exp\left(\theta^{\dagger}c(A')\right)}$$

This implementation aims to reduce the number of times that the support needs to be computed. Models included here use more than a single array, and thus allow the function to recycle support sets as needed. For example, if we are looking at directed graphs all of the same size and without vertex level features, i.e. a model that only counts edges, triangles, etc. then the support needs to be fully computed only once.

Template Parameters

Array_Type	Class of BArray object.
Data_Counter_Type	Any type.
Data_Rule_Type	Any type.

Definition at line 102 of file model-bones.hpp.

7.12.2 Constructor & Destructor Documentation

7.12.2.1 Model() [1/3]

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type >
Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::Model [inline]
```

Definition at line 7 of file model-meat.hpp.

7.12.2.2 Model() [2/3]

Definition at line 27 of file model-meat.hpp.

7.12.2.3 Model() [3/3]

Definition at line 50 of file model-meat.hpp.

7.12.2.4 ∼Model()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_↔
Rule_Type = bool>
Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::~Model ( ) [inline]
```

Definition at line 197 of file model-bones.hpp.

7.12.3 Member Function Documentation

7.12.3.1 add_array()

Adds an array to the support of not already included.

Parameters

Array_	array to be added	
force_new	If false, it will use keygen to obtain a double vector and create a hash of it. If the hash has	
	been computed earlier, the support is recycled.	

Returns

The number of the array.

When computing with the powerset, we need to grow the corresponding vectors on the fly

Definition at line 229 of file model-meat.hpp.

7.12.3.2 add_counter() [1/3]

Definition at line 131 of file model-meat.hpp.

7.12.3.3 add counter() [2/3]

Definition at line 140 of file model-meat.hpp.

7.12.3.4 add_counter() [3/3]

Definition at line 150 of file model-meat.hpp.

7.12.3.5 add_rule() [1/3]

Definition at line 182 of file model-meat.hpp.

7.12.3.6 add_rule() [2/3]

Definition at line 191 of file model-meat.hpp.

7.12.3.7 add_rule() [3/3]

Definition at line 201 of file model-meat.hpp.

7.12.3.8 get norm const()

Definition at line 460 of file model-meat.hpp.

7.12.3.9 get_pset()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type > const std::vector< Array_Type > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type > \leftrightarrow ::get_pset ( const uint & i ) [inline]
```

Definition at line 492 of file model-meat.hpp.

7.12.3.10 get_stats()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type > const std::vector< std::vector< double > > * Model< Array_Type, Data_Counter_Type, Data_\leftarrow Rule_Type >::get_stats ( const uint & i ) [inline]
```

Definition at line 505 of file model-meat.hpp.

7.12.3.11 likelihood() [1/3]

Definition at line 346 of file model-meat.hpp.

7.12.3.12 likelihood() [2/3]

Definition at line 386 of file model-meat.hpp.

7.12.3.13 likelihood() [3/3]

Definition at line 313 of file model-meat.hpp.

7.12.3.14 likelihood_total()

Definition at line 420 of file model-meat.hpp.

7.12.3.15 nterms()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type >
uint Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::nterms [inline], [noexcept]
```

Definition at line 546 of file model-meat.hpp.

7.12.3.16 operator=()

Definition at line 80 of file model-meat.hpp.

7.12.3.17 print_stats()

Definition at line 517 of file model-meat.hpp.

7.12.3.18 sample() [1/2]

7.12.3.19 sample() [2/2]

Definition at line 553 of file model-meat.hpp.

7.12.3.20 set_counters()

Definition at line 169 of file model-meat.hpp.

7.12.3.21 set keygen()

Definition at line 123 of file model-meat.hpp.

7.12.3.22 set_rengine()

Definition at line 113 of file model-bones.hpp.

7.12.3.23 set_rules()

Definition at line 218 of file model-meat.hpp.

7.12.3.24 set seed()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftrightarrow Rule_Type = bool> void Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::set_seed ( unsigned int s ) [inline]
```

Definition at line 123 of file model-bones.hpp.

7.12.3.25 size()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type >
uint Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::size [inline], [noexcept]
```

Definition at line 536 of file model-meat.hpp.

7.12.3.26 size_unique()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type >
uint Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::size_unique [inline], [noexcept]
```

Definition at line 541 of file model-meat.hpp.

7.12.3.27 store_psets()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type >
void Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::store_psets [inline], [noexcept]
```

Definition at line 115 of file model-meat.hpp.

7.12.4 Member Data Documentation

7.12.4.1 array frequency

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
std::vector< uint > Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::array_frequency
```

Definition at line 161 of file model-bones.hpp.

7.12.4.2 arrays2support

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_↔ Rule_Type = bool> std::vector< uint > Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::arrays2support
```

Definition at line 162 of file model-bones.hpp.

7.12.4.3 counter_fun

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
StatsCounter<Array_Type, Data_Counter_Type> Model< Array_Type, Data_Counter_Type, Data_Rule_←
Type >::counter_fun
```

Definition at line 179 of file model-bones.hpp.

7.12.4.4 counters

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
Counters<Array_Type, Data_Counter_Type> Model< Array_Type, Data_Counter_Type, Data_Rule_Type
>::counters
```

Definition at line 176 of file model-bones.hpp.

7.12.4.5 delete_rengine

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::delete_rengine = false
```

Definition at line 112 of file model-bones.hpp.

7.12.4.6 first_calc_done

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
std::vector< bool > Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::first_calc_done
```

Definition at line 185 of file model-bones.hpp.

7.12.4.7 keygen

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarrow Rule_Type = bool> std::function<std::vector<double>const Array_Type &)> Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::keygen = nullptr
```

Function to extract features of the array to be hash.

Definition at line 189 of file model-bones.hpp.

7.12.4.8 keys2support

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
MapVec_type< double, uint > Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::keys2support
```

Map of types of arrays to support sets.

This is of the same length as the vector stats.

Definition at line 169 of file model-bones.hpp.

7.12.4.9 n_arrays_per_stats

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
std::vector< uint > Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::n_arrays_per_←
stats
```

Definition at line 138 of file model-bones.hpp.

7.12.4.10 normalizing_constants

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←> Rule_Type = bool> std::vector< double > Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::normalizing_←> constants
```

Definition at line 184 of file model-bones.hpp.

7.12.4.11 params_last

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
std::vector< std::vector<double> > Model< Array_Type, Data_Counter_Type, Data_Rule_Type >←
::params_last
```

Vector of the previously used parameters.

Definition at line 183 of file model-bones.hpp.

7.12.4.12 pset_arrays

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←> Rule_Type = bool> std::vector< std::vector< Array_Type > > Model< Array_Type, Data_Counter_Type, Data_Rule_←> Type >::pset_arrays
```

Definition at line 147 of file model-bones.hpp.

7.12.4.13 pset probs

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←> Rule_Type = bool> std::vector< std::vector<double> > Model< Array_Type, Data_Counter_Type, Data_Rule_Type > ←> ::pset_probs
```

Definition at line 149 of file model-bones.hpp.

7.12.4.14 pset stats

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
std::vector< std::vector< std::vector<double> > Model< Array_Type, Data_Counter_Type,
Data_Rule_Type >::pset_stats
```

Definition at line 148 of file model-bones.hpp.

7.12.4.15 rengine

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
std::mt19937* Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::rengine = nullptr
```

Definition at line 111 of file model-bones.hpp.

7.12.4.16 rules

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
Rules<Array_Type, Data_Rule_Type> Model< Array_Type, Data_Counter_Type, Data_Rule_Type> ←
::rules
```

Definition at line 177 of file model-bones.hpp.

7.12.4.17 stats

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
std::vector< Counts_type > Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::stats
```

Definition at line 137 of file model-bones.hpp.

7.12.4.18 support_fun

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
Support<Array_Type, Data_Counter_Type, Data_Rule_Type> Model< Array_Type, Data_Counter_Type,
Data_Rule_Type >::support_fun
```

Definition at line 178 of file model-bones.hpp.

7.12.4.19 target_stats

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←> Rule_Type = bool> std::vector< std::vector< double > > Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::target_stats
```

Definition at line 160 of file model-bones.hpp.

7.12.4.20 with_pset

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_↔
Rule_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::with_pset = false
```

Definition at line 146 of file model-bones.hpp.

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

- include/barry/model-bones.hpp
- include/barry/model-meat.hpp

7.13 NetCounterData Class Reference

Data class used to store arbitrary uint or double vectors.

```
#include <network.hpp>
```

Public Member Functions

- NetCounterData ()
- NetCounterData (const std::vector< uint > indices_, const std::vector< double > numbers_)
- ∼NetCounterData ()

Public Attributes

- std::vector< uint > indices
- std::vector< double > numbers

7.13.1 Detailed Description

Data class used to store arbitrary uint or double vectors.

Definition at line 61 of file network.hpp.

7.13.2 Constructor & Destructor Documentation

7.13.2.1 NetCounterData() [1/2]

```
NetCounterData::NetCounterData ( ) [inline]
```

Definition at line 67 of file network.hpp.

7.13.2.2 NetCounterData() [2/2]

Definition at line 68 of file network.hpp.

7.13.2.3 ∼NetCounterData()

```
NetCounterData::~NetCounterData ( ) [inline]
```

Definition at line 73 of file network.hpp.

7.13.3 Member Data Documentation

7.13.3.1 indices

```
std::vector< uint > NetCounterData::indices
```

Definition at line 64 of file network.hpp.

7.13.3.2 numbers

```
std::vector< double > NetCounterData::numbers
```

Definition at line 65 of file network.hpp.

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

• include/barry/counters/network.hpp

7.14 NetworkData Class Reference

Data class for Networks.

```
#include <network.hpp>
```

Public Member Functions

- NetworkData ()
- NetworkData (std::vector< double > vertex_attr_, bool directed_=true)

Constructor using a single attribute.

- NetworkData (std::vector< std::vector< double > > vertex_attr_, bool directed_=true)
 Constructor using multiple attributes.
- ∼NetworkData ()

Public Attributes

- bool directed = true
- $std::vector < std::vector < double > > vertex_attr$

7.14.1 Detailed Description

Data class for Networks.

Details on the available counters for NetworkData can be found in the Network counters section.

This holds information about whether the graph is directed or not, and, if defined, vectors of node (vertex) attributes (vertex_attr).

Definition at line 24 of file network.hpp.

7.14.2 Constructor & Destructor Documentation

7.14.2.1 NetworkData() [1/3]

```
NetworkData::NetworkData ( ) [inline]
```

Definition at line 30 of file network.hpp.

7.14.2.2 NetworkData() [2/3]

Constructor using a single attribute.

Parameters

vertex_← attr_	Double vector of length equal to the number of vertices in the data.
directed_	When true the graph as treated as directed.

Definition at line 38 of file network.hpp.

7.14.2.3 NetworkData() [3/3]

Constructor using multiple attributes.

Parameters

vertex_←	Vector of double vectors. The size equals to the number of attributes to be created. Each	
attr_	individual vector should be of length equal to the number of vertices.	
directed_	When true the graph as treated as directed.	

Definition at line 50 of file network.hpp.

7.14.2.4 ~NetworkData()

NetworkData::~NetworkData () [inline]

Definition at line 56 of file network.hpp.

7.14.3 Member Data Documentation

7.14.3.1 directed

bool NetworkData::directed = true

Definition at line 27 of file network.hpp.

7.14.3.2 vertex_attr

std::vector< std::vector< double > > NetworkData::vertex_attr

Definition at line 28 of file network.hpp.

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

• include/barry/counters/network.hpp

7.15 Node Class Reference

A single node for the model.

#include <geese-node-bones.hpp>

Collaboration diagram for Node:



7.15 Node Class Reference 97

Public Member Functions

- ∼Node ()
- int get_parent () const
- bool is_leaf () const noexcept

Construct a new Node object

- Node ()
- Node (unsigned int id_, unsigned int ord_, bool duplication_)
- Node (unsigned int id_, unsigned int ord_, std::vector< unsigned int > annotations_, bool duplication_)
- Node (Node &&x) noexcept
- Node (const Node &x)

Public Attributes

· unsigned int id

Id of the node (as specified in the input)

· unsigned int ord

Order in which the node was created.

- phylocounters::PhyloArray array
- std::vector< unsigned int > annotations

Observed annotations (only defined for Geese)

- bool duplication
- std::vector< phylocounters::PhyloArray > arrays = {}

Arrays given all possible states.

• Node * parent = nullptr

Parent node.

std::vector < Node * > offspring = {}

Offspring nodes.

• std::vector< unsigned int > narray = {}

ID of the array in the model.

- bool visited = false
- std::vector< double > subtree_prob

Induced subtree probabilities.

std::vector< double > probability

The probability of observing each state.

7.15.1 Detailed Description

A single node for the model.

Each node contains all the information to compute the conditional probability of the pruning algorithm at that node.

Definition at line 11 of file geese-node-bones.hpp.

7.15.2 Constructor & Destructor Documentation

7.15.2.1 Node() [1/5]

```
Node::Node ( ) [inline]
```

Definition at line 36 of file geese-node-bones.hpp.

7.15.2.2 Node() [2/5]

```
Node::Node (
          unsigned int id_,
          unsigned int ord_,
          bool duplication_ ) [inline]
```

Definition at line 55 of file geese-node-bones.hpp.

7.15.2.3 Node() [3/5]

```
Node::Node (
          unsigned int id_,
          unsigned int ord_,
          std::vector< unsigned int > annotations_,
          bool duplication_ ) [inline]
```

Definition at line 61 of file geese-node-bones.hpp.

7.15.2.4 Node() [4/5]

Definition at line 68 of file geese-node-bones.hpp.

7.15.2.5 Node() [5/5]

Definition at line 82 of file geese-node-bones.hpp.

7.15 Node Class Reference 99

7.15.2.6 ∼Node()

```
Node::~Node ( ) [inline]
```

Definition at line 47 of file geese-node-bones.hpp.

7.15.3 Member Function Documentation

7.15.3.1 get_parent()

```
int Node::get_parent ( ) const [inline]
```

Definition at line 96 of file geese-node-bones.hpp.

7.15.3.2 is_leaf()

```
bool Node::is_leaf ( ) const [inline], [noexcept]
```

Definition at line 103 of file geese-node-bones.hpp.

7.15.4 Member Data Documentation

7.15.4.1 annotations

```
std::vector< unsigned int > Node::annotations
```

Observed annotations (only defined for Geese)

Definition at line 18 of file geese-node-bones.hpp.

7.15.4.2 array

phylocounters::PhyloArray Node::array

Definition at line 17 of file geese-node-bones.hpp.

7.15.4.3 arrays

```
std::vector< phylocounters::PhyloArray > Node::arrays = {}
```

Arrays given all possible states.

Definition at line 21 of file geese-node-bones.hpp.

7.15.4.4 duplication

```
bool Node::duplication
```

Definition at line 19 of file geese-node-bones.hpp.

7.15.4.5 id

```
unsigned int Node::id
```

Id of the node (as specified in the input)

Definition at line 14 of file geese-node-bones.hpp.

7.15.4.6 narray

```
std::vector< unsigned int > Node::narray = {}
```

ID of the array in the model.

Definition at line 24 of file geese-node-bones.hpp.

7.15.4.7 offspring

```
std::vector< Node* > Node::offspring = {}
```

Offspring nodes.

Definition at line 23 of file geese-node-bones.hpp.

7.15 Node Class Reference 101

7.15.4.8 ord

unsigned int Node::ord

Order in which the node was created.

Definition at line 15 of file geese-node-bones.hpp.

7.15.4.9 parent

```
Node* Node::parent = nullptr
```

Parent node.

Definition at line 22 of file geese-node-bones.hpp.

7.15.4.10 probability

```
std::vector< double > Node::probability
```

The probability of observing each state.

Definition at line 28 of file geese-node-bones.hpp.

7.15.4.11 subtree prob

```
std::vector< double > Node::subtree_prob
```

Induced subtree probabilities.

Definition at line 27 of file geese-node-bones.hpp.

7.15.4.12 visited

```
bool Node::visited = false
```

Definition at line 25 of file geese-node-bones.hpp.

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

• include/barry/models/geese/geese-node-bones.hpp

7.16 NodeData Class Reference

Data definition for the PhyloArray class.

```
#include <phylo.hpp>
```

Public Member Functions

- · NodeData ()
- NodeData (const std::vector< double > &blengths_, const std::vector< bool > &states_, bool duplication
 —=true)
- ∼NodeData ()

Public Attributes

- std::vector< double > blengths
- std::vector< bool > states
- bool duplication = true

7.16.1 Detailed Description

Data definition for the PhyloArray class.

Details about the available counters for PhyloArray objects can be found in the Phylo counters section.

This holds basic information about a given node.

Definition at line 23 of file phylo.hpp.

7.16.2 Constructor & Destructor Documentation

7.16.2.1 NodeData() [1/2]

```
NodeData::NodeData ( ) [inline]
```

Definition at line 41 of file phylo.hpp.

7.16.2.2 NodeData() [2/2]

Definition at line 43 of file phylo.hpp.

7.16.2.3 ~NodeData()

```
NodeData::~NodeData ( ) [inline]
```

Definition at line 49 of file phylo.hpp.

7.16.3 Member Data Documentation

7.16.3.1 blengths

```
std::vector< double > NodeData::blengths
```

Branch length.

Definition at line 29 of file phylo.hpp.

7.16.3.2 duplication

```
bool NodeData::duplication = true
```

Definition at line 39 of file phylo.hpp.

7.16.3.3 states

```
\verb|std::vector<|bool|> \verb|NodeData::states|\\
```

State of the parent node.

Definition at line 34 of file phylo.hpp.

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

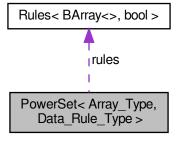
• include/barry/counters/phylo.hpp

7.17 PowerSet< Array_Type, Data_Rule_Type > Class Template Reference

Powerset of a binary array.

#include <powerset-bones.hpp>

Collaboration diagram for PowerSet < Array_Type, Data_Rule_Type >:



Public Member Functions

- void init support ()
- void calc ()
- void reset (uint N_, uint M_)

Construct and destroy a PowerSet object

- PowerSet ()
- PowerSet (uint N_, uint M_)
- PowerSet (const Array_Type & array)
- ∼PowerSet ()

Wrappers for the <tt>Rules</tt> member.

These will add rules to the model, which are shared by the support and the actual counter function.

- void add rule (Rule < Array Type, Data Rule Type > &rule)
- void add_rule (Rule < Array_Type, Data_Rule_Type > *rule)
- void add_rule (Rule_fun_type< Array_Type, Data_Rule_Type > count_fun_, Data_Rule_Type *data_
 =nullptr, bool delete_data_=false)

Getter functions

- const std::vector< Array_Type > * get_data_ptr () const
- std::vector< Array_Type > get_data () const
- std::vector< Array_Type >::iterator begin ()
- std::vector< Array_Type >::iterator end ()
- std::size_t size () const noexcept
- const Array_Type & operator[] (const unsigned int &i) const

Public Attributes

- Array_Type EmptyArray
- std::vector< Array_Type > data
- Rules< Array_Type, Data_Rule_Type > * rules
- uint N
- uint M
- bool rules_deleted = false
- std::vector< std::pair< uint, uint >> coordinates_free
- std::vector< std::pair< uint, uint >> coordinates locked

7.17.1 Detailed Description

```
template<typename Array_Type = BArray<>>, typename Data_Rule_Type = bool> class PowerSet< Array_Type, Data_Rule_Type >
```

Powerset of a binary array.

Template Parameters

Array_Type	
Data_Rule_Type	

Definition at line 17 of file powerset-bones.hpp.

7.17.2 Constructor & Destructor Documentation

7.17.2.1 PowerSet() [1/3]

```
template<typename Array_Type = BArray<>>, typename Data_Rule_Type = bool>
PowerSet< Array_Type, Data_Rule_Type >::PowerSet ( ) [inline]
```

Definition at line 39 of file powerset-bones.hpp.

7.17.2.2 PowerSet() [2/3]

Definition at line 41 of file powerset-bones.hpp.

7.17.2.3 PowerSet() [3/3]

Definition at line 7 of file powerset-meat.hpp.

7.17.2.4 ∼PowerSet()

```
template<typename Array_Type , typename Data_Rule_Type >
PowerSet< Array_Type, Data_Rule_Type >::~PowerSet [inline]
```

Definition at line 15 of file powerset-meat.hpp.

7.17.3 Member Function Documentation

7.17.3.1 add_rule() [1/3]

Definition at line 113 of file powerset-meat.hpp.

7.17.3.2 add_rule() [2/3]

Definition at line 122 of file powerset-meat.hpp.

7.17.3.3 add_rule() [3/3]

```
template<typename Array_Type , typename Data_Rule_Type >
void PowerSet< Array_Type, Data_Rule_Type >::add_rule (
    Rule_fun_type< Array_Type, Data_Rule_Type > count_fun_,
    Data_Rule_Type * data_ = nullptr,
    bool delete_data_ = false ) [inline]
```

Definition at line 132 of file powerset-meat.hpp.

7.17.3.4 begin()

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< Array_Type >::iterator PowerSet< Array_Type, Data_Rule_Type >::begin ( ) [inline]
```

Definition at line 73 of file powerset-bones.hpp.

7.17.3.5 calc()

```
template<typename Array_Type , typename Data_Rule_Type >
void PowerSet< Array_Type, Data_Rule_Type >::calc [inline]
```

Definition at line 88 of file powerset-meat.hpp.

7.17.3.6 end()

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< Array_Type >::iterator PowerSet< Array_Type, Data_Rule_Type >::end ( ) [inline]
```

Definition at line 74 of file powerset-bones.hpp.

7.17.3.7 get_data()

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< Array_Type > PowerSet< Array_Type, Data_Rule_Type >::get_data ( ) const [inline]
```

Definition at line 72 of file powerset-bones.hpp.

7.17.3.8 get_data_ptr()

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
const std::vector< Array_Type >* PowerSet< Array_Type, Data_Rule_Type >::get_data_ptr ()
const [inline]
```

Definition at line 71 of file powerset-bones.hpp.

7.17.3.9 init_support()

```
template<typename Array_Type , typename Data_Rule_Type >
void PowerSet< Array_Type, Data_Rule_Type >::init_support [inline]
```

Definition at line 21 of file powerset-meat.hpp.

7.17.3.10 operator[]()

Definition at line 76 of file powerset-bones.hpp.

7.17.3.11 reset()

Definition at line 101 of file powerset-meat.hpp.

7.17.3.12 size()

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::size_t PowerSet< Array_Type, Data_Rule_Type >::size ( ) const [inline], [noexcept]
```

Definition at line 75 of file powerset-bones.hpp.

7.17.4 Member Data Documentation

7.17.4.1 coordinates_free

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< std::pair<uint,uint> > PowerSet< Array_Type, Data_Rule_Type >::coordinates_free
```

Definition at line 31 of file powerset-bones.hpp.

7.17.4.2 coordinates_locked

template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< std::pair<uint,uint> > PowerSet< Array_Type, Data_Rule_Type >::coordinates_← locked

Definition at line 32 of file powerset-bones.hpp.

7.17.4.3 data

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< Array_Type > PowerSet< Array_Type, Data_Rule_Type >::data
```

Definition at line 24 of file powerset-bones.hpp.

7.17.4.4 EmptyArray

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
Array_Type PowerSet< Array_Type, Data_Rule_Type >::EmptyArray
```

Definition at line 23 of file powerset-bones.hpp.

7.17.4.5 M

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
uint PowerSet< Array_Type, Data_Rule_Type >::M
```

Definition at line 27 of file powerset-bones.hpp.

7.17.4.6 N

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
uint PowerSet< Array_Type, Data_Rule_Type >::N
```

Definition at line 27 of file powerset-bones.hpp.

7.17.4.7 rules

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
Rules<Array_Type, Data_Rule_Type>* PowerSet< Array_Type, Data_Rule_Type >::rules
```

Definition at line 25 of file powerset-bones.hpp.

7.17.4.8 rules_deleted

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
bool PowerSet< Array_Type, Data_Rule_Type >::rules_deleted = false
```

Definition at line 28 of file powerset-bones.hpp.

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

- · include/barry/powerset-bones.hpp
- include/barry/powerset-meat.hpp

7.18 Rule < Array_Type, Data_Type > Class Template Reference

Rule for determining if a cell should be included in a sequence.

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

Public Member Functions

- ∼Rule ()
- bool locked (const Array_Type &a, uint i, uint j)

Construct a new Rule object

Construct a new Rule object

Parameters

fun_	A function of type Rule_fun_type.
dat_	Data pointer to be passed to fun_
delete_←	When true, the Rule destructor will delete the pointer, if defined.
dat_	

- Rule ()
- Rule (Rule_fun_type< Array_Type, Data_Type > fun_, Data_Type *dat_=nullptr, bool delete_dat_=false)

7.18.1 Detailed Description

```
template<typename Array_Type = BArray<>>, typename Data_Type = bool> class Rule< Array_Type, Data_Type>
```

Rule for determining if a cell should be included in a sequence.

Rules can be used together with Support and PowerSet to determine which cells should be included when enumerating all possible realizations of a binary array.

Template Parameters

Array_Type	An object of class BArray.
Data_Type	Any type.

Definition at line 23 of file rules-bones.hpp.

7.18.2 Constructor & Destructor Documentation

7.18.2.1 Rule() [1/2]

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Rule< Array_Type, Data_Type >::Rule ( ) [inline]
```

Definition at line 42 of file rules-bones.hpp.

7.18.2.2 Rule() [2/2]

Definition at line 43 of file rules-bones.hpp.

7.18.2.3 ∼Rule()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Rule< Array_Type, Data_Type >::~Rule ( ) [inline]
```

Definition at line 50 of file rules-bones.hpp.

7.18.3 Member Function Documentation

7.18.3.1 locked()

Definition at line 63 of file rules-meat.hpp.

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

- include/barry/rules-bones.hpp
- include/barry/rules-meat.hpp

7.19 Rules < Array_Type, Data_Type > Class Template Reference

Vector of objects of class Rule.

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

Public Member Functions

- Rules ()
- Rules (const Rules < Array_Type, Data_Type > &rules_)
- Rules< Array_Type, Data_Type > operator= (const Rules< Array_Type, Data_Type > &rules_)
- ∼Rules ()
- uint size () const noexcept
- bool locked (const Array_Type &a, uint i, uint j)

Check whether a given cell is free or locked.

- void clear ()
- void get_seq (const Array_Type &a, std::vector< std::pair< uint, uint >> *free, std::vector< std::pair< uint, uint >> *locked=nullptr)

Computes the sequence of free and locked cells in an BArray.

Rule adding

Parameters

rule

```
    void add_rule (Rule < Array_Type, Data_Type > &rule)
```

```
    void add_rule (Rule < Array_Type, Data_Type > *rule)
```

void add_rule (Rule_fun_type< Array_Type, Data_Type > rule_, Data_Type *data_=nullptr, bool delete
 __data_=false)

7.19.1 Detailed Description

```
template<typename Array_Type, typename Data_Type> class Rules< Array_Type, Data_Type>
```

Vector of objects of class Rule.

Template Parameters

Array_Type	An object of class BArray
Data_Type	Any type.

Definition at line 67 of file rules-bones.hpp.

7.19.2 Constructor & Destructor Documentation

7.19.2.1 Rules() [1/2]

```
template<typename Array_Type , typename Data_Type >
Rules< Array_Type, Data_Type >::Rules ( ) [inline]
```

Definition at line 74 of file rules-bones.hpp.

7.19.2.2 Rules() [2/2]

Definition at line 10 of file rules-meat.hpp.

7.19.2.3 ∼Rules()

```
template<typename Array_Type , typename Data_Type >
Rules< Array_Type, Data_Type >::~Rules ( ) [inline]
```

Definition at line 79 of file rules-bones.hpp.

7.19.3 Member Function Documentation

7.19.3.1 add_rule() [1/3]

Definition at line 68 of file rules-meat.hpp.

7.19.3.2 add_rule() [2/3]

Definition at line 79 of file rules-meat.hpp.

7.19.3.3 add_rule() [3/3]

Definition at line 89 of file rules-meat.hpp.

7.19.3.4 clear()

```
template<typename Array_Type , typename Data_Type >
void Rules< Array_Type, Data_Type >::clear [inline]
```

Definition at line 127 of file rules-meat.hpp.

7.19.3.5 get_seq()

Computes the sequence of free and locked cells in an BArray.

Parameters

а	An object of class BArray.	
free	Pointer to a vector of pairs (i, j) listing the free cells.	
locked	(optional) Pointer to a vector of pairs (i, j) listing the locked cells.	

Returns

Nothing.

Definition at line 139 of file rules-meat.hpp.

7.19.3.6 locked()

Check whether a given cell is free or locked.

Parameters

а	A BArray object
i	row position
j	col position

Returns

true If the cell is locked false If the cell is free

Definition at line 111 of file rules-meat.hpp.

7.19.3.7 operator=()

Definition at line 35 of file rules-meat.hpp.

7.19.3.8 size()

```
template<typename Array_Type , typename Data_Type >
uint Rules< Array_Type, Data_Type >::size ( ) const [inline], [noexcept]
```

Definition at line 84 of file rules-bones.hpp.

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

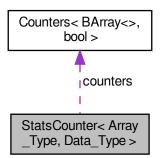
- include/barry/rules-bones.hpp
- include/barry/rules-meat.hpp

7.20 StatsCounter< Array_Type, Data_Type > Class Template Reference

Count stats for a single Array.

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

Collaboration diagram for StatsCounter< Array_Type, Data_Type >:



Public Member Functions

• StatsCounter (const Array_Type *Array_)

Creator of a StatsCounter

StatsCounter ()

Can be created without setting the array.

- ∼StatsCounter ()
- void reset_array (const Array_Type *Array_)

Changes the reference array for the counting.

- void add_counter (Counter< Array_Type, Data_Type > *f_)
- void add_counter (Counter< Array_Type, Data_Type > f_)
- void set_counters (Counters< Array_Type, Data_Type > *counters_)
- void count_init (uint i, uint j)

Counter functions This function recurses through the entries of Array and at each step of adding a new cell it uses the functions to list the statistics.

- void count_current (uint i, uint j)
- std::vector< double > count_all ()

Public Attributes

- const Array_Type * Array
- Array_Type EmptyArray
- std::vector< double > current stats
- $\bullet \ \ Counters < Array_Type, \ Data_Type > * \ counters$
- bool counter_deleted = false

7.20.1 Detailed Description

```
\label{template} $$ \ensuremath{\sf template}$$ $$ $$ \ensuremath{\sf template}$$ $$ $$ \ensuremath{\sf Type}$ = BArray<>, typename Data_Type = bool> class StatsCounter< Array_Type, Data_Type>
```

Count stats for a single Array.

Users can a list of functions that can be used with this. The baseline set of arguments is a pointer to a binary array and a dataset to add the counts to.

Definition at line 16 of file statscounter-bones.hpp.

7.20.2 Constructor & Destructor Documentation

7.20.2.1 StatsCounter() [1/2]

Creator of a StatsCounter

Parameters

Array⊷	A const pointer to a BArray.
_	

Definition at line 34 of file statscounter-bones.hpp.

7.20.2.2 StatsCounter() [2/2]

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
StatsCounter< Array_Type, Data_Type >::StatsCounter ( ) [inline]
```

Can be created without setting the array.

Definition at line 49 of file statscounter-bones.hpp.

7.20.2.3 ~StatsCounter()

```
template<typename Array_Type , typename Data_Type >
StatsCounter< Array_Type, Data_Type >::~StatsCounter [inline]
```

Definition at line 7 of file statscounter-meat.hpp.

7.20.3 Member Function Documentation

7.20.3.1 add_counter() [1/2]

Definition at line 25 of file statscounter-meat.hpp.

7.20.3.2 add counter() [2/2]

Definition at line 35 of file statscounter-meat.hpp.

7.20.3.3 count_all()

```
template<typename Array_Type , typename Data_Type >
std::vector< double > StatsCounter< Array_Type, Data_Type >::count_all [inline]
```

Definition at line 99 of file statscounter-meat.hpp.

7.20.3.4 count_current()

Definition at line 81 of file statscounter-meat.hpp.

7.20.3.5 count_init()

Counter functions This function recurses through the entries of Array and at each step of adding a new cell it uses the functions to list the statistics.

Definition at line 61 of file statscounter-meat.hpp.

7.20.3.6 reset array()

Changes the reference array for the counting.

Parameters

Array←	A pointer to an array of class Array_Type.

Definition at line 14 of file statscounter-meat.hpp.

7.20.3.7 set_counters()

Definition at line 46 of file statscounter-meat.hpp.

7.20.4 Member Data Documentation

7.20.4.1 Array

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
const Array_Type* StatsCounter< Array_Type, Data_Type >::Array
```

Definition at line 21 of file statscounter-bones.hpp.

7.20.4.2 counter_deleted

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
bool StatsCounter< Array_Type, Data_Type >::counter_deleted = false
```

Definition at line 27 of file statscounter-bones.hpp.

7.20.4.3 counters

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Counters<Array_Type,Data_Type>* StatsCounter< Array_Type, Data_Type >::counters
```

Definition at line 26 of file statscounter-bones.hpp.

7.20.4.4 current_stats

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
std::vector< double > StatsCounter< Array_Type, Data_Type >::current_stats
```

Definition at line 23 of file statscounter-bones.hpp.

7.20.4.5 EmptyArray

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Array_Type StatsCounter< Array_Type, Data_Type >::EmptyArray
```

Definition at line 22 of file statscounter-bones.hpp.

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

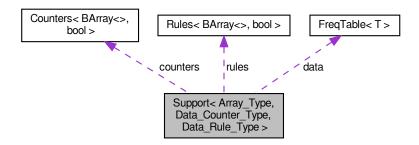
- include/barry/statscounter-bones.hpp
- include/barry/statscounter-meat.hpp

7.21 Support< Array_Type, Data_Counter_Type, Data_Rule_Type > Class Template Reference

Compute the support of sufficient statistics.

#include <support-bones.hpp>

Collaboration diagram for Support < Array Type, Data Counter Type, Data Rule Type >:



Public Member Functions

- Support (const Array_Type &Array_)
 - Constructor passing a reference Array.
- Support (uint N_, uint M_)

Constructor specifying the dimensions of the array (empty).

- Support ()
- ∼Support ()
- void init_support (std::vector< Array_Type > *array_bank=nullptr, std::vector< std::vector< double > >
 *stats_bank=nullptr)
- void calc (std::vector< Array_Type > *array_bank=nullptr, std::vector< std::vector< double >> *stats_\cup bank=nullptr, unsigned int max_num_elements_=0u)

Computes the entire support.

- Counts_type get_counts () const
- const MapVec_type * get_counts_ptr () const
- void print () const

Resets the support calculator

If needed, the counters of a support object can be reused.

Parameters

Aı	rray←	New array over which the support will be computed.
l _		

- void reset array ()
- void reset_array (const Array_Type &Array_)

Manage counters

Parameters

f_	A counter to be added.
counters←	A vector of counters to be added.
_	

- void add_counter (Counter< Array_Type, Data_Counter_Type > *f_)
- void add_counter (Counter< Array_Type, Data_Counter_Type > f_)
- void set_counters (Counters < Array_Type, Data_Counter_Type > *counters_)

Manage rules

Parameters

f_	A rule to be added.
counters←	A vector of rules to be added.
_	

- void add_rule (Rule < Array_Type, Data_Rule_Type > *f_)
- void add_rule (Rule < Array_Type, Data_Rule_Type > f_)
- void set_rules (Rules < Array_Type, Data_Rule_Type > *rules_)

Public Attributes

- Array_Type EmptyArray
 - Reference array to generate the support.
- · FreqTable data
- Counters
 Array_Type, Data_Counter_Type > * counters
- Rules
 Array_Type, Data_Rule_Type > * rules
- uint N
- uint M
- bool counter_deleted = false
- bool rules_deleted = false
- uint max_num_elements = BARRY_MAX_NUM_ELEMENTS
- std::vector< double > current_stats
- std::vector< std::pair< uint, uint >> coordinates_free
- $\bullet \ \ \mathsf{std} :: \mathsf{vector} < \mathsf{std} :: \mathsf{pair} < \mathsf{uint}, \ \mathsf{uint} > > \mathsf{coordinates_locked} \\$
- std::vector< std::vector< double > > change_stats

7.21.1 Detailed Description

template < typename Array_Type = BArray <>>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool > class Support < Array_Type, Data_Counter_Type, Data_Rule_Type >

Compute the support of sufficient statistics.

Given an array and a set of counters, this object iterates throughout the support set of the Array while at the same time computing the support of the sufficient statitics.

Definition at line 24 of file support-bones.hpp.

7.21.2 Constructor & Destructor Documentation

7.21.2.1 Support() [1/3]

Constructor passing a reference Array.

Definition at line 56 of file support-bones.hpp.

7.21.2.2 Support() [2/3]

Constructor specifying the dimensions of the array (empty).

Definition at line 64 of file support-bones.hpp.

7.21.2.3 Support() [3/3]

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
Support< Array_Type, Data_Counter_Type, Data_Rule_Type >::Support ( ) [inline]
```

Definition at line 70 of file support-bones.hpp.

7.21.2.4 ~Support()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_↔
Rule_Type = bool>
Support< Array_Type, Data_Counter_Type, Data_Rule_Type >::~Support ( ) [inline]
```

Definition at line 76 of file support-bones.hpp.

7.21.3 Member Function Documentation

7.21.3.1 add_counter() [1/2]

Definition at line 178 of file support-meat.hpp.

7.21.3.2 add_counter() [2/2]

Definition at line 188 of file support-meat.hpp.

7.21.3.3 add_rule() [1/2]

Definition at line 215 of file support-meat.hpp.

7.21.3.4 add_rule() [2/2]

Definition at line 225 of file support-meat.hpp.

7.21.3.5 calc()

Computes the entire support.

Not to be used by the user. Sets the starting point in the array (column-major).

Parameters

array_bank	If specified, the counter will add to the vector each possible state of the array, as it counts.
stats_bank	If specified, the counter will add to the vector each possible set of statistics, as it counts.

Definition at line 152 of file support-meat.hpp.

7.21.3.6 get_counts()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type >
Counts_type Support< Array_Type, Data_Counter_Type, Data_Rule_Type >::get_counts [inline]
```

Definition at line 252 of file support-meat.hpp.

7.21.3.7 get_counts_ptr()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type >
const MapVec_type * Support< Array_Type, Data_Counter_Type, Data_Rule_Type >::get_counts_ptr
[inline]
```

Definition at line 259 of file support-meat.hpp.

7.21.3.8 init_support()

Definition at line 7 of file support-meat.hpp.

7.21.3.9 print()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type >
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type >::print [inline]
```

Definition at line 266 of file support-meat.hpp.

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7.21.3.10 reset_array() [1/2]

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type >
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type >::reset_array [inline]
```

Definition at line 67 of file support-meat.hpp.

7.21.3.11 reset_array() [2/2]

Definition at line 74 of file support-meat.hpp.

7.21.3.12 set_counters()

Definition at line 198 of file support-meat.hpp.

7.21.3.13 set_rules()

Definition at line 235 of file support-meat.hpp.

7.21.4 Member Data Documentation

7.21.4.1 change_stats

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarray_Rule_Type = bool>
std::vector< std::vector< double > > Support< Array_Type, Data_Counter_Type, Data_Rule_Type
>::change_stats
```

Definition at line 52 of file support-bones.hpp.

7.21.4.2 coordinates_free

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←> Rule_Type = bool> std::vector< std::pair<uint,uint> > Support< Array_Type, Data_Counter_Type, Data_Rule_Type >::coordinates_free
```

Definition at line 50 of file support-bones.hpp.

7.21.4.3 coordinates locked

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
std::vector< std::pair<uint,uint> > Support< Array_Type, Data_Counter_Type, Data_Rule_Type
>::coordinates_locked
```

Definition at line 51 of file support-bones.hpp.

7.21.4.4 counter deleted

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type >::counter_deleted = false
```

Definition at line 44 of file support-bones.hpp.

7.21.4.5 counters

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
Counters<Array_Type, Data_Counter_Type>* Support< Array_Type, Data_Counter_Type, Data_Rule_←
Type >::counters
```

Definition at line 40 of file support-bones.hpp.

7.21.4.6 current_stats

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\top
Rule_Type = bool>
std::vector< double > Support< Array_Type, Data_Counter_Type, Data_Rule_Type >::current_stats
```

Definition at line 49 of file support-bones.hpp.

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7.21.4.7 data

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
FreqTable Support< Array_Type, Data_Counter_Type, Data_Rule_Type >::data
```

Definition at line 39 of file support-bones.hpp.

7.21.4.8 EmptyArray

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
Array_Type Support< Array_Type, Data_Counter_Type, Data_Rule_Type >::EmptyArray
```

Reference array to generate the support.

Definition at line 38 of file support-bones.hpp.

7.21.4.9 M

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
uint Support< Array_Type, Data_Counter_Type, Data_Rule_Type >::M
```

Definition at line 43 of file support-bones.hpp.

7.21.4.10 max_num_elements

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_↔
Rule_Type = bool>
uint Support< Array_Type, Data_Counter_Type, Data_Rule_Type >::max_num_elements = BARRY_MAX_NUM_ELEMENTS
```

Definition at line 46 of file support-bones.hpp.

7.21.4.11 N

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
uint Support< Array_Type, Data_Counter_Type, Data_Rule_Type >::N
```

Definition at line 43 of file support-bones.hpp.

7.21.4.12 rules

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
Rules<Array_Type, Data_Rule_Type>* Support< Array_Type, Data_Counter_Type, Data_Rule_Type >←
::rules
```

Definition at line 41 of file support-bones.hpp.

7.21.4.13 rules deleted

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type >::rules_deleted = false
```

Definition at line 45 of file support-bones.hpp.

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

- include/barry/support-bones.hpp
- include/barry/support-meat.hpp

7.22 vecHasher< T > Struct Template Reference

```
#include <typedefs.hpp>
```

Public Member Functions

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

7.22.1 Detailed Description

```
template<typename T> struct vecHasher< T>
```

Definition at line 97 of file typedefs.hpp.

7.22.2 Member Function Documentation

7.22.2.1 operator()()

Definition at line 98 of file typedefs.hpp.

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

include/barry/typedefs.hpp

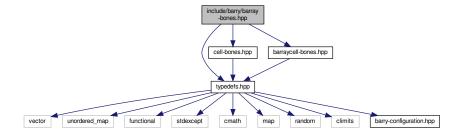
130 Class Documentation

Chapter 8

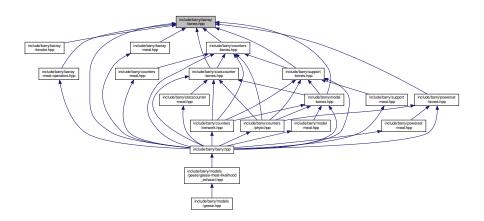
File Documentation

8.1 include/barry/barray-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "cell-bones.hpp"
#include "barraycell-bones.hpp"
Include dependency graph for barray-bones.hpp:
```



This graph shows which files directly or indirectly include this file:



Classes

class BArray < Cell_Type, Data_Type >
 Baseline class for binary arrays.

Macros

• #define BARRAY_BONES_HPP 1

8.1.1 Macro Definition Documentation

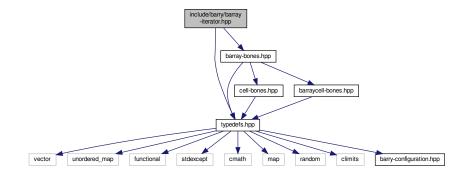
8.1.1.1 BARRAY_BONES_HPP

```
#define BARRAY_BONES_HPP 1
```

Definition at line 8 of file barray-bones.hpp.

8.2 include/barry/barray-iterator.hpp File Reference

```
#include "typedefs.hpp"
#include "barray-bones.hpp"
Include dependency graph for barray-iterator.hpp:
```



Classes

 $\bullet \ \, {\sf class\ ConstBArrayRowlter} < {\sf Cell_Type,\ Data_Type} >$

Macros

#define BARRAY_ITERATOR_HPP 1

8.2.1 Macro Definition Documentation

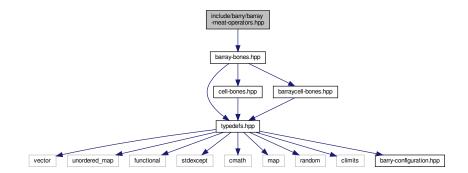
8.2.1.1 BARRAY_ITERATOR_HPP

#define BARRAY_ITERATOR_HPP 1

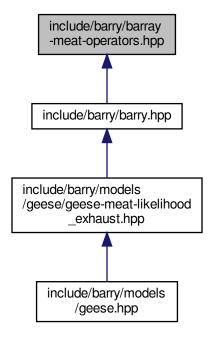
Definition at line 7 of file barray-iterator.hpp.

8.3 include/barry/barray-meat-operators.hpp File Reference

#include "barray-bones.hpp"
Include dependency graph for barray-meat-operators.hpp:



This graph shows which files directly or indirectly include this file:



Functions

template<typename Cell_Type , typename Data_Type >
 void checkdim_ (const BArray< Cell_Type, Data_Type > &lhs, const BArray< Cell_Type, Data_Type > &rhs)

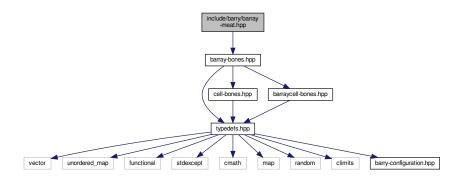
8.3.1 Function Documentation

8.3.1.1 checkdim_()

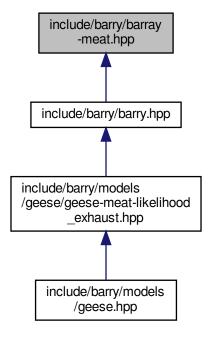
Definition at line 8 of file barray-meat-operators.hpp.

8.4 include/barry/barray-meat.hpp File Reference

#include "barray-bones.hpp"
Include dependency graph for barray-meat.hpp:



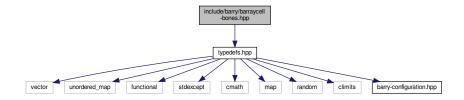
This graph shows which files directly or indirectly include this file:



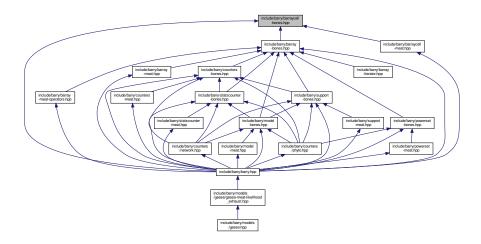
8.5 include/barry/barraycell-bones.hpp File Reference

#include "typedefs.hpp"

Include dependency graph for barraycell-bones.hpp:



This graph shows which files directly or indirectly include this file:

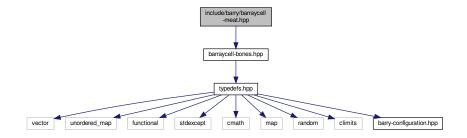


Classes

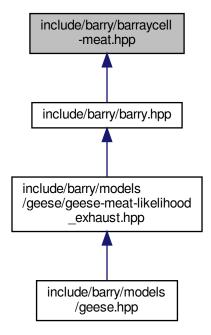
- class BArrayCell< Cell_Type, Data_Type>
- class BArrayCell_const< Cell_Type, Data_Type >

8.6 include/barry/barraycell-meat.hpp File Reference

#include "barraycell-bones.hpp"
Include dependency graph for barraycell-meat.hpp:

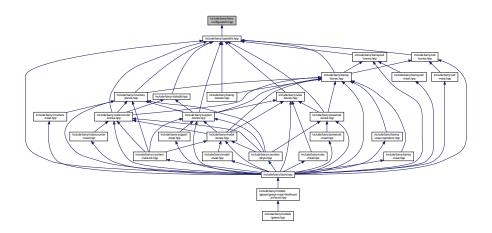


This graph shows which files directly or indirectly include this file:



8.7 include/barry/barry-configuration.hpp File Reference

This graph shows which files directly or indirectly include this file:



Macros

• #define BARRY_MAX_NUM_ELEMENTS static_cast< unsigned int >(UINT_MAX/2u)

Configuration MACROS

These are mostly related to performance. The definitions follow:

- BARRY_USE_UNORDERED_MAP If specified, then barry is compiled using std::unordered_map. Otherwise it will use std::map for the arrays.
- BARRY_USE_SAFE_EXP When specified, it will multiply all likelihoods in Model by (1/-100)/(1/-100) so that numerical overflows are avoided.
- BARRY_CHECK_FINITE When specified, it will introduce a macro

```
• #define BARRY_SAFE_EXP -100.0
```

- #define BARRY_ISFINITE(a)
- #define BARRY CHECK SUPPORT(x, maxs)
- template<typename Ta , typename Tb >
 using Map = std::map< Ta, Tb >

8.7.1 Macro Definition Documentation

8.7.1.1 BARRY_CHECK_SUPPORT

Definition at line 46 of file barry-configuration.hpp.

8.7.1.2 BARRY ISFINITE

```
#define BARRY_ISFINITE(
```

Definition at line 39 of file barry-configuration.hpp.

8.7.1.3 BARRY_MAX_NUM_ELEMENTS

```
\texttt{\#define BARRY\_MAX\_NUM\_ELEMENTS static\_cast} < \text{unsigned int } > \texttt{(UINT\_MAX/2u)}
```

Definition at line 5 of file barry-configuration.hpp.

8.7.1.4 BARRY_SAFE_EXP

```
#define BARRY_SAFE_EXP -100.0
```

Definition at line 32 of file barry-configuration.hpp.

8.7.2 Typedef Documentation

8.7.2.1 Map

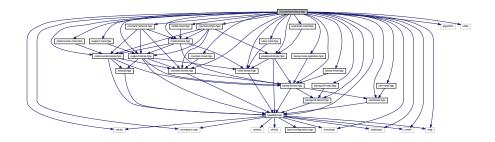
```
template<typename Ta , typename Tb >
using Map = std::map<Ta,Tb>
```

Definition at line 26 of file barry-configuration.hpp.

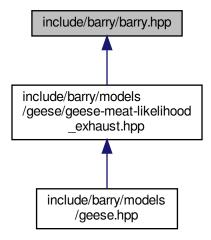
8.8 include/barry/barry.hpp File Reference

```
#include <vector>
#include <unordered_map>
#include <functional>
#include <stdexcept>
#include <cmath>
#include <map>
#include <algorithm>
#include <utility>
#include "typedefs.hpp"
#include "cell-bones.hpp"
#include "cell-meat.hpp"
#include "barray-bones.hpp"
#include "barraycell-bones.hpp"
#include "barray-meat.hpp"
#include "barraycell-meat.hpp"
#include "barray-meat-operators.hpp"
#include "counters-bones.hpp"
#include "counters-meat.hpp"
#include "statscounter-bones.hpp"
#include "statscounter-meat.hpp"
#include "support-bones.hpp"
#include "support-meat.hpp"
#include "powerset-bones.hpp"
#include "powerset-meat.hpp"
#include "model-bones.hpp"
#include "model-meat.hpp"
#include "rules-bones.hpp"
#include "rules-meat.hpp"
#include "counters/network.hpp"
```

#include "counters/phylo.hpp"
Include dependency graph for barry.hpp:



This graph shows which files directly or indirectly include this file:



Namespaces

barry

barry: Your go-to motif accountant

· barry::counters

Tree class and Treelterator class.

- barry::counters::network
- · barry::counters::phylo

Macros

- #define COUNTER_FUNCTION(a)
- #define COUNTER_LAMBDA(a)
- #define RULE_FUNCTION(a)
- #define RULE_LAMBDA(a)

8.8.1 Macro Definition Documentation

8.8.1.1 COUNTER_FUNCTION

```
#define COUNTER_FUNCTION( a )
```

Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
inline double (a) (const Array_Type & Array, uint i, uint j, Data_Type * data) \
```

Definition at line 64 of file barry.hpp.

8.8.1.2 COUNTER_LAMBDA

```
\#define COUNTER_LAMBDA( a )
```

Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
Counter_fun_type<Array_Type, Data_Type> a = \
[](const Array_Type & Array, uint i, uint j, Data_Type * data)
```

Definition at line 67 of file barry.hpp.

8.8.1.3 RULE FUNCTION

```
#define RULE_FUNCTION( a )
```

Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
inline bool (a) (const Array_Type & Array, uint i, uint j, Data_Type * data)\
```

Definition at line 71 of file barry.hpp.

8.8.1.4 RULE_LAMBDA

```
#define RULE_LAMBDA( a )
```

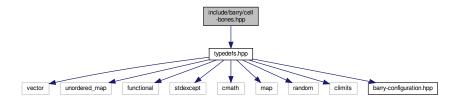
Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
Rule_fun_type<Array_Type, Data_Type> a = \
[](const Array_Type & Array, uint i, uint j, Data_Type * data)
```

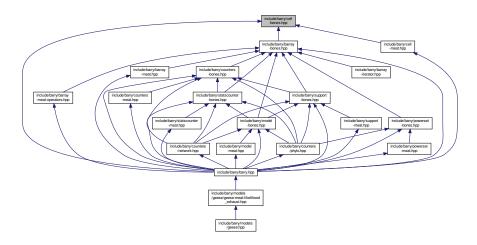
Definition at line 74 of file barry.hpp.

8.9 include/barry/cell-bones.hpp File Reference

#include "typedefs.hpp"
Include dependency graph for cell-bones.hpp:



This graph shows which files directly or indirectly include this file:



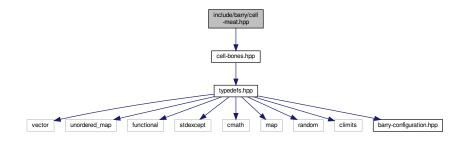
Classes

• class Cell< Cell_Type >

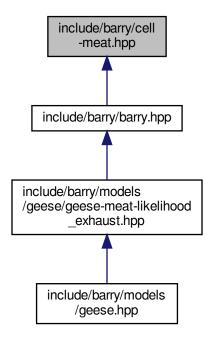
Entries in BArray. For now, it only has two members:

8.10 include/barry/cell-meat.hpp File Reference

#include "cell-bones.hpp"
Include dependency graph for cell-meat.hpp:



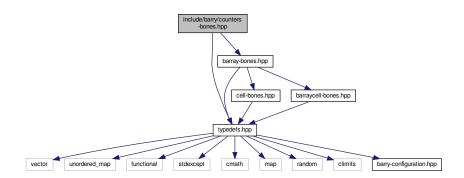
This graph shows which files directly or indirectly include this file:



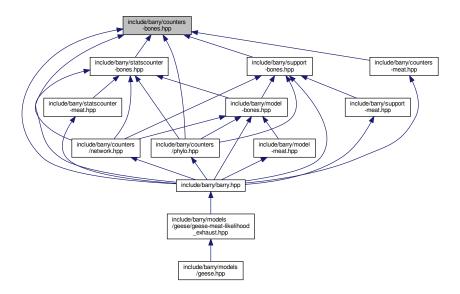
8.11 include/barry/col-bones.hpp File Reference

8.12 include/barry/counters-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "barray-bones.hpp"
Include dependency graph for counters-bones.hpp:
```



This graph shows which files directly or indirectly include this file:



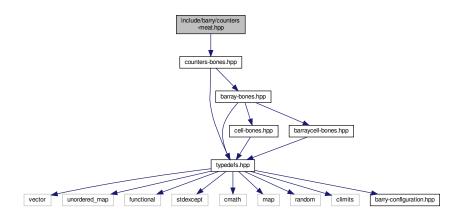
Classes

- class Counter< Array_Type, Data_Type >
 - A counter function based on change statistics.
- $\bullet \ \ {\it class Counters}{< Array_Type, \, Data_Type} >$

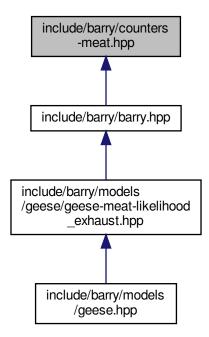
Vector of counters.

8.13 include/barry/counters-meat.hpp File Reference

#include "counters-bones.hpp"
Include dependency graph for counters-meat.hpp:



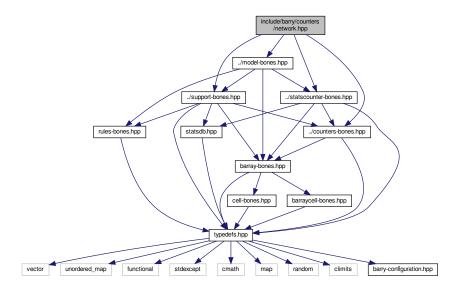
This graph shows which files directly or indirectly include this file:



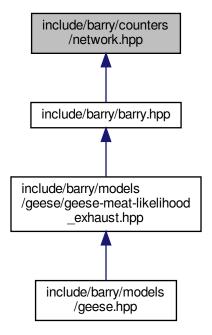
8.14 include/barry/counters/network.hpp File Reference

```
#include "../counters-bones.hpp"
#include "../support-bones.hpp"
#include "../statscounter-bones.hpp"
#include "../model-bones.hpp"
```

Include dependency graph for network.hpp:



This graph shows which files directly or indirectly include this file:



Classes

class NetworkData

Data class for Networks.

· class NetCounterData

Data class used to store arbitrary uint or double vectors.

Macros

- #define NET_C_DATA_IDX(i) (data->indices[i])
- #define NET_C_DATA_NUM(i) (data->numbers[i])

Macros for defining counters

- #define NETWORK_COUNTER(a)
- #define NETWORK_COUNTER_LAMBDA(a)

Macros for defining rules

- #define NETWORK RULE(a)
- #define NETWORK_RULE_LAMBDA(a)

Typedefs

Convenient typedefs for network objects.

- typedef BArray< double, NetworkData > Network
- typedef Counter
 NetCounterData > NetCounter
- typedef Counters < Network, NetCounterData > NetCounters
- typedef Support < Network, NetCounterData > NetSupport
- $\bullet \ \ typedef \ StatsCounter < Network, \ NetCounter Data > NetStatsCounter \\$
- typedef Model < Network, NetCounterData > NetModel
- typedef Rule < Network, bool > NetRule
- typedef Rules < Network, bool > NetRules

Functions

void counter_edges (NetCounters *counters)

Number of edges.

• void counter_isolates (NetCounters *counters)

Number of isolated vertices.

void counter_mutual (NetCounters *counters)

Number of mutual ties.

- void counter_istar2 (NetCounters *counters)
- void counter_ostar2 (NetCounters *counters)
- void counter ttriads (NetCounters *counters)
- void counter_ctriads (NetCounters *counters)
- void counter_density (NetCounters *counters)
- void counter_idegree15 (NetCounters *counters)
- void counter_odegree15 (NetCounters *counters)
- void counter_absdiff (NetCounters *counters, uint attr_id, double alpha=1.0)

Sum of absolute attribute difference between ego and alter.

- void counter_diff (NetCounters *counters, uint attr_id, double alpha=1.0, double tail_head=true)

 Sum of attribute difference between ego and alter to pow(alpha)
- NETWORK_COUNTER (init_single_attr)
- void counter_nodeicov (NetCounters *counters, uint attr_id)
- void counter nodeocov (NetCounters *counters, uint attr id)
- void counter_nodecov (NetCounters *counters, uint attr_id)
- · void counter_nodematch (NetCounters *counters, uint attr_id)
- void counter_idegree (NetCounters *counters, std::vector< uint > d)

Counts number of vertices with a given in-degree.

void counter_odegree (NetCounters *counters, std::vector< uint > d)

Counts number of vertices with a given out-degree.

void counter_degree (NetCounters *counters, std::vector< uint > d)

Counts number of vertices with a given out-degree.

Rules for network models

Parameters

```
rules | A pointer to a NetRules object (Rules < Network, bool > ).
```

void rules_zerodiag (NetRules *rules)
 Number of edges.

8.14.1 Macro Definition Documentation

8.14.1.1 **NET_C_DATA_IDX**

Definition at line 79 of file network.hpp.

8.14.1.2 NET_C_DATA_NUM

Definition at line 80 of file network.hpp.

8.14.1.3 NETWORK_COUNTER

Function for definition of a network counter function

Definition at line 101 of file network.hpp.

8.14.1.4 NETWORK_COUNTER_LAMBDA

Lambda function for definition of a network counter function

Definition at line 104 of file network.hpp.

8.14.1.5 NETWORK_RULE

Function for definition of a network counter function

Definition at line 113 of file network.hpp.

8.14.1.6 NETWORK RULE LAMBDA

Lambda function for definition of a network counter function

Definition at line 116 of file network.hpp.

8.14.2 Typedef Documentation

8.14.2.1 NetCounter

typedef Counter<Network, NetCounterData > NetCounter

Definition at line 88 of file network.hpp.

8.14.2.2 NetCounters

typedef Counters< Network, NetCounterData> NetCounters

Definition at line 89 of file network.hpp.

8.14.2.3 NetModel

typedef Model<Network, NetCounterData> NetModel

Definition at line 92 of file network.hpp.

8.14.2.4 NetRule

typedef Rule<Network,bool> NetRule

Definition at line 93 of file network.hpp.

8.14.2.5 NetRules

typedef Rules<Network,bool> NetRules

Definition at line 94 of file network.hpp.

8.14.2.6 NetStatsCounter

```
typedef StatsCounter<Network, NetCounterData> NetStatsCounter
```

Definition at line 91 of file network.hpp.

8.14.2.7 NetSupport

```
typedef Support<Network, NetCounterData > NetSupport
```

Definition at line 90 of file network.hpp.

8.14.2.8 Network

```
typedef BArray<double, NetworkData> Network
```

Definition at line 87 of file network.hpp.

8.14.3 Function Documentation

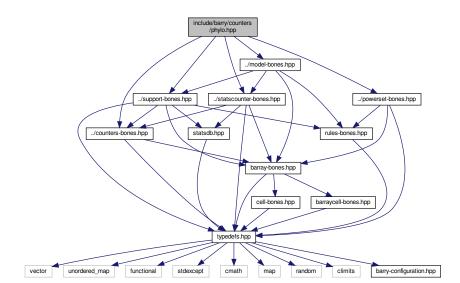
8.14.3.1 rules_zerodiag()

Number of edges.

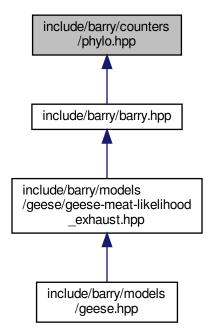
Definition at line 742 of file network.hpp.

8.15 include/barry/counters/phylo.hpp File Reference

```
#include "../counters-bones.hpp"
#include "../support-bones.hpp"
#include "../statscounter-bones.hpp"
#include "../model-bones.hpp"
#include "../powerset-bones.hpp"
Include dependency graph for phylo.hpp:
```



This graph shows which files directly or indirectly include this file:



Classes

· class NodeData

Data definition for the PhyloArray class.

Macros

- #define PHYLO COUNTER(a)
 - Extension of a simple counter.
- #define PHYLO COUNTER LAMBDA(a)
- #define PHYLO_CHECK_MISSING()

Typedefs

Convenient typedefs for Node objects.

- typedef BArray< uint, NodeData > PhyloArray
- typedef Counter< PhyloArray, PhyloCounterData > PhyloCounter
- typedef Counters< PhyloArray, PhyloCounterData > PhyloCounters
- typedef Rule< PhyloArray, PhyloRuleData > PhyloRule
- typedef Rules < PhyloArray, PhyloRuleData > PhyloRules
- typedef Support < PhyloArray, PhyloCounterData, PhyloRuleData > PhyloSupport
- typedef StatsCounter< PhyloArray, PhyloCounterData > PhyloStatsCounter
- typedef Model < PhyloArray, PhyloCounterData, PhyloRuleData > PhyloModel
- typedef PowerSet
 PhyloArray, PhyloRuleData > PhyloPowerSet

Functions

• void counter_overall_gains (PhyloCounters *counters, bool duplication=true)

Overall functional gains.

void counter_gains (PhyloCounters *counters, std::vector< uint > nfun, bool duplication=true)

Functional gains for a specific function (nfun).

void counter_gains_k_offspring (PhyloCounters *counters, std::vector< uint > nfun, uint k=1u, bool duplication=true)

Functional gains for a specific function (nfun).

void counter genes changing (PhyloCounters *counters, bool duplication=true)

Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)

• void counter_overall_loss (PhyloCounters *counters, bool duplication=true)

Overall functional loss.

void counter_maxfuns (PhyloCounters *counters, uint lb, uint ub, bool duplication=true)

Cap the number of functions per gene.

void counter_loss (PhyloCounters *counters, std::vector< uint > nfun, bool duplication=true)

Total count of losses for an specific function.

void counter_overall_changes (PhyloCounters *counters, bool duplication=true)

Total number of changes. Use this statistic to account for "preservation".

• void counter subfun (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)

Total count of Sub-functionalization events.

• void counter_cogain (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)

Co-evolution (joint gain or loss)

void counter_longest (PhyloCounters *counters)

Longest branch mutates (either by gain or by loss)

• void counter_neofun (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)

Total number of neofunctionalization events.

• void counter neofun a2b (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)

Total number of neofunctionalization events.

void counter_co_opt (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)
 Function co-opting.

- #define PHYLO_C_DATA_IDX(i) (data.operator[](i))
- typedef std::vector< uint > PhyloCounterData
- typedef std::vector< std::pair< uint, uint > > PhyloRuleData

8.15.1 Macro Definition Documentation

8.15.1.1 PHYLO_C_DATA_IDX

Definition at line 56 of file phylo.hpp.

8.15.1.2 PHYLO_CHECK_MISSING

```
#define PHYLO_CHECK_MISSING( )

Value:
    if (Array.data == nullptr) \
        throw std::logic_error("The array data is nullptr."); \
        if (data == nullptr) \
        throw std::logic_error("The counter data is nullptr.")
```

Definition at line 91 of file phylo.hpp.

8.15.1.3 PHYLO_COUNTER

Extension of a simple counter.

It allows specifying extra arguments, in particular, the corresponding sets of rows to which this statistic may be relevant. This could be important in the case of, for example, counting correlation type statistics between function 1 and 2, and between function 1 and 3.

Definition at line 85 of file phylo.hpp.

8.15.1.4 PHYLO_COUNTER_LAMBDA

Definition at line 88 of file phylo.hpp.

8.15.2 Typedef Documentation

8.15.2.1 PhyloArray

```
typedef BArray<uint, NodeData> PhyloArray
```

Definition at line 63 of file phylo.hpp.

8.15.2.2 PhyloCounter

typedef Counter<PhyloArray, PhyloCounterData > PhyloCounter

Definition at line 64 of file phylo.hpp.

8.15.2.3 PhyloCounterData

typedef std::vector< uint > PhyloCounterData

Definition at line 53 of file phylo.hpp.

8.15.2.4 PhyloCounters

typedef Counters< PhyloArray, PhyloCounterData> PhyloCounters

Definition at line 65 of file phylo.hpp.

8.15.2.5 PhyloModel

typedef Model<PhyloArray, PhyloCounterData, PhyloRuleData> PhyloModel

Definition at line 70 of file phylo.hpp.

8.15.2.6 PhyloPowerSet

typedef PowerSet<PhyloArray, PhyloRuleData> PhyloPowerSet

Definition at line 71 of file phylo.hpp.

8.15.2.7 PhyloRule

typedef Rule<PhyloArray,PhyloRuleData> PhyloRule

Definition at line 66 of file phylo.hpp.

8.15.2.8 PhyloRuleData

```
typedef std::vector< std::pair< uint, uint > > PhyloRuleData
```

Definition at line 54 of file phylo.hpp.

8.15.2.9 PhyloRules

```
typedef Rules<PhyloArray,PhyloRuleData> PhyloRules
```

Definition at line 67 of file phylo.hpp.

8.15.2.10 PhyloStatsCounter

```
typedef StatsCounter<PhyloArray, PhyloCounterData> PhyloStatsCounter
```

Definition at line 69 of file phylo.hpp.

8.15.2.11 PhyloSupport

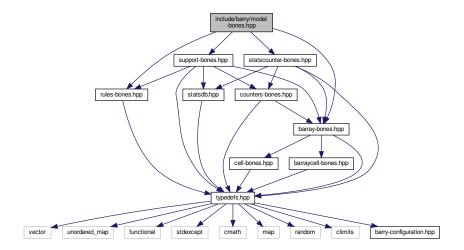
typedef Support < PhyloArray, PhyloCounterData, PhyloRuleData > PhyloSupport

Definition at line 68 of file phylo.hpp.

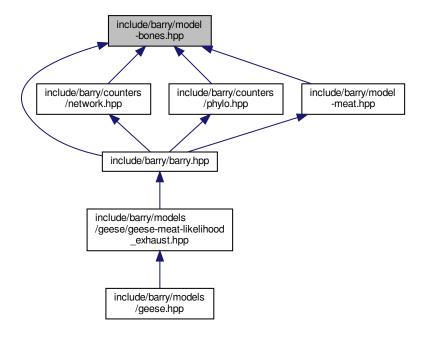
8.16 include/barry/model-bones.hpp File Reference

```
#include "barray-bones.hpp"
#include "support-bones.hpp"
#include "statscounter-bones.hpp"
#include "rules-bones.hpp"
```

Include dependency graph for model-bones.hpp:



This graph shows which files directly or indirectly include this file:



Classes

class Model < Array_Type, Data_Counter_Type, Data_Rule_Type >

General framework for discrete exponential models. This class allows generating discrete exponential models in the form of a linear exponential model:

Functions

- double update normalizing constant (const std::vector< double > ¶ms, const Counts type &support)
- double likelihood_ (const std::vector< double > &target_stats, const std::vector< double > ¶ms, const double normalizing_constant, bool log_=false)
- template<typename Array_Type >
 std::vector< double > keygen_default (const Array_Type &Array_)
 Array Hasher class (used for computing support)

8.16.1 Function Documentation

8.16.1.1 keygen_default()

Array Hasher class (used for computing support)

Definition at line 69 of file model-bones.hpp.

8.16.1.2 likelihood_()

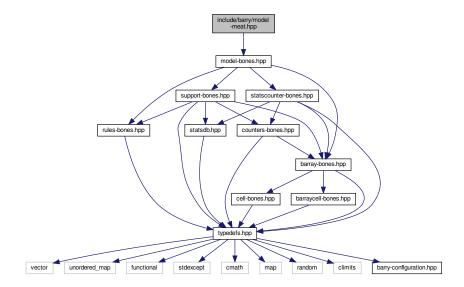
Definition at line 40 of file model-bones.hpp.

8.16.1.3 update_normalizing_constant()

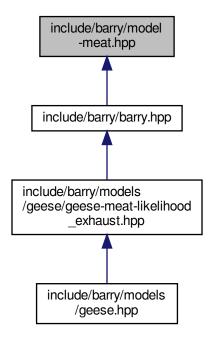
Definition at line 16 of file model-bones.hpp.

8.17 include/barry/model-meat.hpp File Reference

```
#include "model-bones.hpp"
Include dependency graph for model-meat.hpp:
```

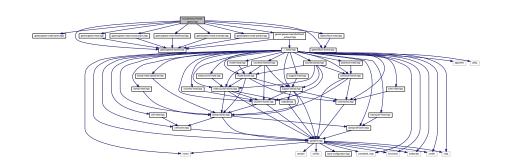


This graph shows which files directly or indirectly include this file:



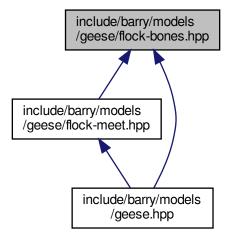
8.18 include/barry/models/geese.hpp File Reference

```
#include "geese/geese-node-bones.hpp"
#include "geese/geese-bones.hpp"
#include "geese/geese-meat.hpp"
#include "geese/geese-meat-constructors.hpp"
#include "geese/geese-meat-likelihood.hpp"
#include "geese/geese-meat-likelihood_exhaust.hpp"
#include "geese/geese-meat-simulate.hpp"
#include "geese/geese-meat-predict.hpp"
#include "geese/flock-bones.hpp"
#include "geese/flock-meet.hpp"
Include dependency graph for geese.hpp:
```



8.19 include/barry/models/geese/flock-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



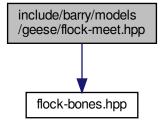
Classes

class Flock

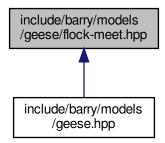
A Flock is a group of Geese.

8.20 include/barry/models/geese/flock-meet.hpp File Reference

#include "flock-bones.hpp"
Include dependency graph for flock-meet.hpp:

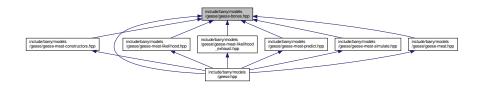


This graph shows which files directly or indirectly include this file:



8.21 include/barry/models/geese/geese-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



Classes

• class Geese

Annotated Phylo Model.

Macros

• #define INITIALIZED()

Functions

- template<typename Ta , typename Tb > std::vector< Ta > vector_caster (const std::vector< Tb > &x)
- RULE_FUNCTION (rule_empty_free)
- std::vector< double > keygen_full (const phylocounters::PhyloArray &array)
- bool vec_diff (const std::vector< unsigned int > &s, const std::vector< unsigned int > &a)

8.21.1 Macro Definition Documentation

8.21.1.1 INITIALIZED

```
#define INITIALIZED( )

Value:
    if (!this->initialized) \
        throw std::logic_error("The model has not been initialized yet.");
```

Definition at line 18 of file geese-bones.hpp.

8.21.2 Function Documentation

8.21.2.1 keygen_full()

Definition at line 31 of file geese-bones.hpp.

8.21.2.2 RULE_FUNCTION()

```
RULE_FUNCTION (
          rule_empty_free )
```

Definition at line 22 of file geese-bones.hpp.

8.21.2.3 vec_diff()

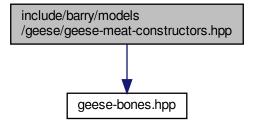
Definition at line 51 of file geese-bones.hpp.

8.21.2.4 vector_caster()

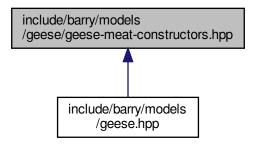
Definition at line 10 of file geese-bones.hpp.

8.22 include/barry/models/geese/geese-meat-constructors.hpp File Reference

```
#include "geese-bones.hpp"
Include dependency graph for geese-meat-constructors.hpp:
```



This graph shows which files directly or indirectly include this file:



Macros

• #define GEESE_MEAT_CONSTRUCTORS_HPP 1

8.22.1 Macro Definition Documentation

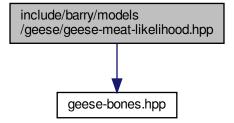
8.22.1.1 GEESE_MEAT_CONSTRUCTORS_HPP

#define GEESE_MEAT_CONSTRUCTORS_HPP 1

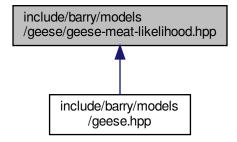
Definition at line 4 of file geese-meat-constructors.hpp.

8.23 include/barry/models/geese/geese-meat-likelihood.hpp File Reference

#include "geese-bones.hpp"
Include dependency graph for geese-meat-likelihood.hpp:

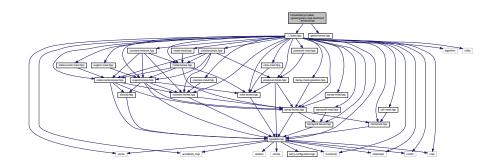


This graph shows which files directly or indirectly include this file:

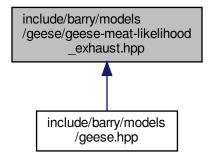


8.24 include/barry/models/geese/geese-meat-likelihood_exhaust.hpp File Reference

#include "../../barry.hpp"
#include "geese-bones.hpp"
Include dependency graph for geese-meat-likelihood_exhaust.hpp:



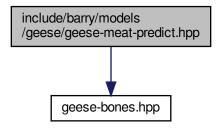
This graph shows which files directly or indirectly include this file:



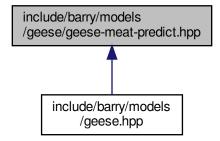
8.25 include/barry/models/geese/geese-meat-predict.hpp File Reference

#include "geese-bones.hpp"

Include dependency graph for geese-meat-predict.hpp:



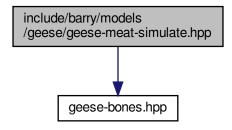
This graph shows which files directly or indirectly include this file:



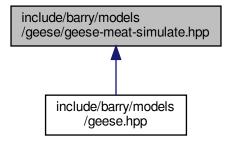
8.26 include/barry/models/geese/geese-meat-simulate.hpp File Reference

#include "geese-bones.hpp"

Include dependency graph for geese-meat-simulate.hpp:

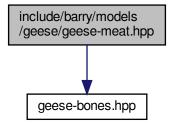


This graph shows which files directly or indirectly include this file:

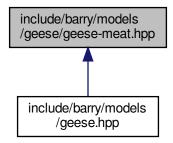


8.27 include/barry/models/geese/geese-meat.hpp File Reference

#include "geese-bones.hpp"
Include dependency graph for geese-meat.hpp:

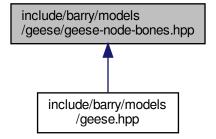


This graph shows which files directly or indirectly include this file:



8.28 include/barry/models/geese/geese-node-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



Classes

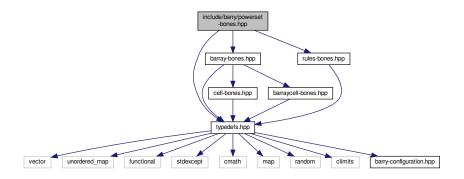
• class Node

A single node for the model.

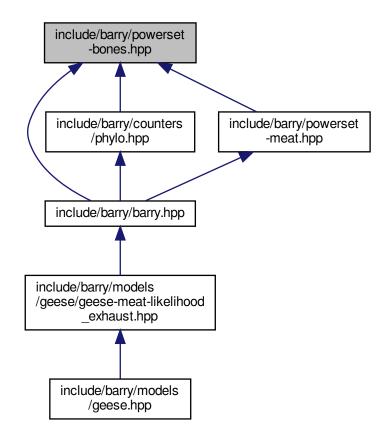
8.29 include/barry/powerset-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "barray-bones.hpp"
```

#include "rules-bones.hpp"
Include dependency graph for powerset-bones.hpp:



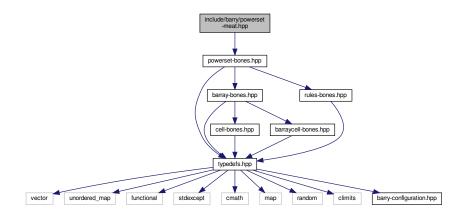
This graph shows which files directly or indirectly include this file:



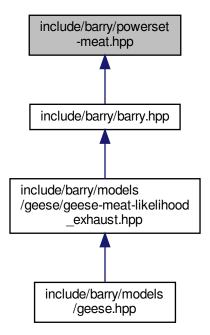
Classes

8.30 include/barry/powerset-meat.hpp File Reference

#include "powerset-bones.hpp"
Include dependency graph for powerset-meat.hpp:



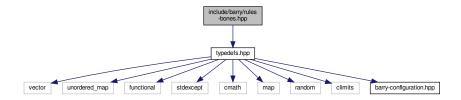
This graph shows which files directly or indirectly include this file:



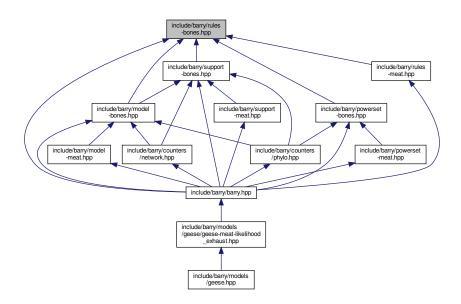
8.31 include/barry/rules-bones.hpp File Reference

#include "typedefs.hpp"

Include dependency graph for rules-bones.hpp:



This graph shows which files directly or indirectly include this file:



Classes

- class Rule < Array_Type, Data_Type >
 - Rule for determining if a cell should be included in a sequence.
- class Rules < Array_Type, Data_Type >

Vector of objects of class Rule.

Functions

template<typename Array_Type , typename Data_Type >
 bool rule_fun_default (const Array_Type *array, uint i, uint j, Data_Type *dat)

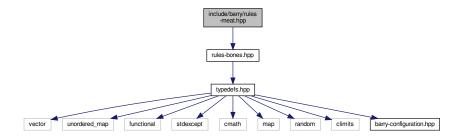
8.31.1 Function Documentation

8.31.1.1 rule_fun_default()

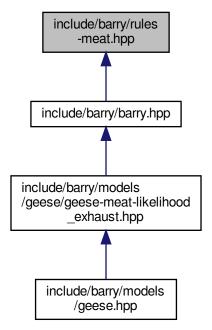
Definition at line 10 of file rules-bones.hpp.

8.32 include/barry/rules-meat.hpp File Reference

#include "rules-bones.hpp"
Include dependency graph for rules-meat.hpp:



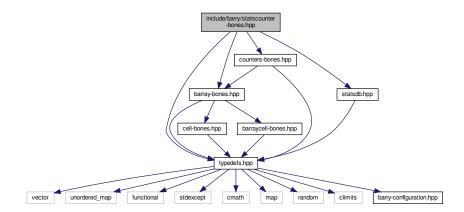
This graph shows which files directly or indirectly include this file:



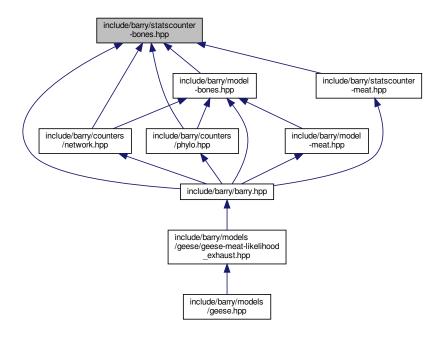
8.33 include/barry/statscounter-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "barray-bones.hpp"
#include "statsdb.hpp"
#include "counters-bones.hpp"
```

Include dependency graph for statscounter-bones.hpp:



This graph shows which files directly or indirectly include this file:

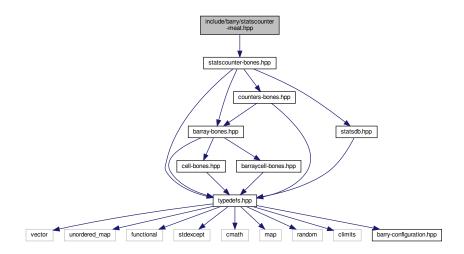


Classes

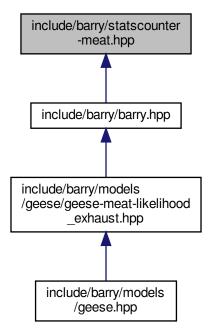
class StatsCounter < Array_Type, Data_Type >
 Count stats for a single Array.

8.34 include/barry/statscounter-meat.hpp File Reference

#include "statscounter-bones.hpp"
Include dependency graph for statscounter-meat.hpp:

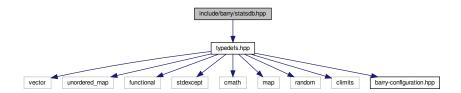


This graph shows which files directly or indirectly include this file:

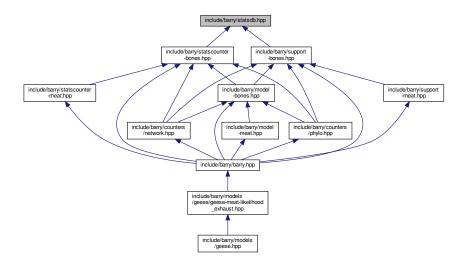


8.35 include/barry/statsdb.hpp File Reference

#include "typedefs.hpp"
Include dependency graph for statsdb.hpp:



This graph shows which files directly or indirectly include this file:



Classes

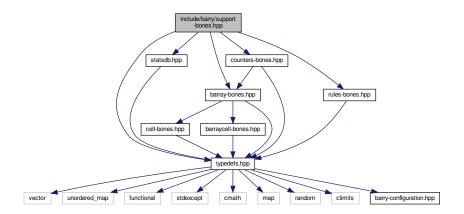
class FreqTable < T >

Database of statistics.

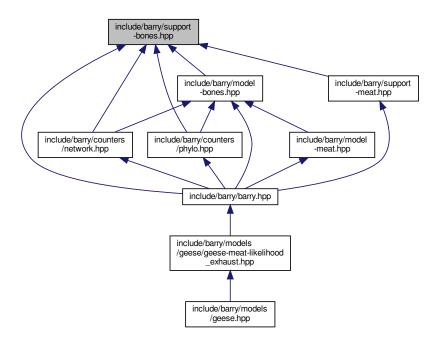
8.36 include/barry/support-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "barray-bones.hpp"
#include "statsdb.hpp"
#include "counters-bones.hpp"
```

#include "rules-bones.hpp"
Include dependency graph for support-bones.hpp:



This graph shows which files directly or indirectly include this file:

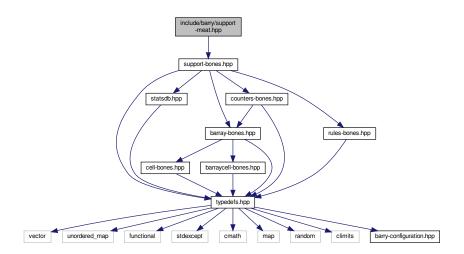


Classes

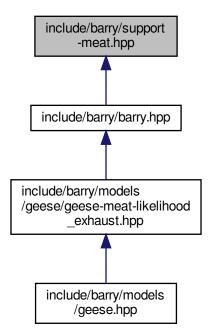
class Support < Array_Type, Data_Counter_Type, Data_Rule_Type >
 Compute the support of sufficient statistics.

8.37 include/barry/support-meat.hpp File Reference

#include "support-bones.hpp"
Include dependency graph for support-meat.hpp:



This graph shows which files directly or indirectly include this file:



Macros

• #define BARRY_SUPPORT_MEAT_HPP 1

8.37.1 Macro Definition Documentation

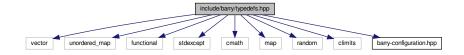
8.37.1.1 BARRY_SUPPORT_MEAT_HPP

```
#define BARRY_SUPPORT_MEAT_HPP 1
```

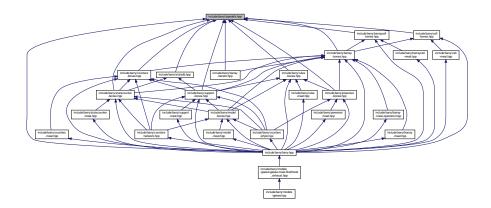
Definition at line 4 of file support-meat.hpp.

8.38 include/barry/typedefs.hpp File Reference

```
#include <vector>
#include <unordered_map>
#include <functional>
#include <stdexcept>
#include <cmath>
#include <map>
#include <random>
#include <climits>
#include "barry-configuration.hpp"
Include dependency graph for typedefs.hpp:
```



This graph shows which files directly or indirectly include this file:



Classes

- class Entries < Cell_Type >
 - A wrapper class to store source, target, val from a BArray object.
- struct vecHasher< T >

Namespaces

CHECK

Integer constants used to specify which cell should be check.

• EXISTS

Integer constants used to specify which cell should be check to exist or not.

Macros

- #define ROW(a) this->el_ij[a]
- #define COL(a) this->el_ji[a]
- #define A_ROW(a) Array.el_ij[a]
- #define A_COL(a) Array.el_ji[a]

Typedefs

- typedef unsigned int uint
- typedef std::vector< std::vector< double >, uint > > Counts type
- template<typename Cell_Type >
 using Row_type = Map< uint, Cell< Cell_Type > >
- template<typename Cell_Type >
 using Col_type = Map< uint, Cell< Cell_Type > * >
- template<typename Ta = double, typename Tb = uint>
 using MapVec_type = std::unordered_map< std::vector< Ta >, Tb, vecHasher< Ta > >
- template<typename Array_Type , typename Data_Type >
 using Counter_fun_type = std::function< double(const Array_Type &, uint, uint, Data_Type *)>
 Counter and rule functions.
- template<typename Array_Type , typename Data_Type >
 using Rule_fun_type = std::function< bool(const Array_Type &, uint, uint, Data_Type *)>

Functions

- template<typename T >
 T vec_inner_prod (const std::vector< T > &a, const std::vector< T > &b)
- template<typename T >
 bool vec_equal (const std::vector< T > &a, const std::vector< T > &b)
 Compares if -a- and -b- are equal.
- $\begin{tabular}{ll} \bullet & template < typename T > \\ & bool vec_equal_approx (const std::vector < T > \&a, const std::vector < T > \&b, double eps=1e-10) \\ \end{tabular}$

Variables

```
const int CHECK::BOTH = -1
const int CHECK::NONE = 0
const int CHECK::ONE = 1
const int CHECK::TWO = 2
const int EXISTS::BOTH = -1
const int EXISTS::NONE = 0
const int EXISTS::ONE = 1
const int EXISTS::TWO = 1
const int EXISTS::UKNOWN = -1
const int EXISTS::AS_ZERO = 0
const int EXISTS::AS_ONE = 1
```

8.38.1 Macro Definition Documentation

8.38.1.1 A_COL

Definition at line 123 of file typedefs.hpp.

8.38.1.2 A_ROW

Definition at line 122 of file typedefs.hpp.

8.38.1.3 COL

Definition at line 23 of file typedefs.hpp.

8.38.1.4 ROW

```
#define ROW( a \ ) \ \  \  \text{this->el_ij[a]}
```

Definition at line 22 of file typedefs.hpp.

8.38.2 Typedef Documentation

8.38.2.1 Col_type

```
template<typename Cell_Type >
using Col_type = Map< uint, Cell<Cell_Type>* >
```

Definition at line 62 of file typedefs.hpp.

8.38.2.2 Counter_fun_type

```
template<typename Array_Type , typename Data_Type >
using Counter_fun_type = std::function<double(const Array_Type &, uint, uint, Data_Type *)>
```

Counter and rule functions.

Parameters

Array_Type	a BArray
unit,uint	Focal cell
Data_Type	Data associated with the function, for example, id of the attribute in the Array.

Returns

```
Counter_fun_type a double (the change statistic)
Rule_fun_type a bool. True if the cell is blocked.
```

Definition at line 136 of file typedefs.hpp.

8.38.2.3 Counts_type

```
typedef std::vector< std::pair< std::vector<double>, uint > > Counts_type
```

Definition at line 55 of file typedefs.hpp.

8.38.2.4 MapVec_type

```
template<typename Ta = double, typename Tb = uint>
using MapVec_type = std::unordered_map< std::vector< Ta >, Tb, vecHasher<Ta> >
```

Definition at line 116 of file typedefs.hpp.

8.38.2.5 Row_type

```
template<typename Cell_Type >
using Row_type = Map< uint, Cell<Cell_Type> >
```

Definition at line 59 of file typedefs.hpp.

8.38.2.6 Rule_fun_type

```
template<typename Array_Type , typename Data_Type >
using Rule_fun_type = std::function<bool(const Array_Type &, uint, uint, Data_Type *)>
```

Definition at line 139 of file typedefs.hpp.

8.38.2.7 uint

```
typedef unsigned int uint
```

Definition at line 19 of file typedefs.hpp.

8.38.3 Function Documentation

8.38.3.1 vec_equal()

Compares if -a- and -b- are equal.

Parameters

```
a,b Two vectors of the same length
```

Returns

true if all elements are equal.

Definition at line 150 of file typedefs.hpp.

8.38.3.2 vec_equal_approx()

Definition at line 168 of file typedefs.hpp.

8.38.3.3 vec_inner_prod()

Definition at line 188 of file typedefs.hpp.

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