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);
```

2 Main Page

```
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.

# **Module Index**

## 2.1 Modules

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Network counters						 											 					10
Phylo counters						 											 					15

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# **Class Index**

## 3.1 Class List

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

BArray < Cell_Type, Data_Type >
Baseline class for binary arrays
BArrayCell < Cell_Type, Data_Type >
BArrayCell_const< Cell_Type, Data_Type >
Cell< Cell_Type >
Entries in BArray. For now, it only has two members:
ConstBArrayRowlter< Cell_Type, Data_Type >
Counter< Array_Type, Data_Type >
A counter function based on change statistics
Counters < Array_Type, Data_Type >
Vector of counters
Entries < Cell_Type >
A wrapper class to store source, target, val from a BArray object 5
Flock
A Flock is a group of Geese
FreqTable < T >
Database of statistics
Geese
Annotated Phylo Model
Model< Array_Type, Data_Counter_Type, Data_Rule_Type >
General framework for discrete exponential models. This class allows generating discrete expo-
nential models in the form of a linear exponential model:
NetCounterData
Data class used to store arbitrary uint or double vectors
NetworkData
Data class for Networks
Node
A single node for the model
NodeData
Data definition for the PhyloArray class 10
PowerSet < Array_Type, Data_Rule_Type >
Powerset of a binary array
Rule < Array_Type, Data_Type >
Rule for determining if a cell should be included in a sequence
Rules< Array_Type, Data_Type >
Vector of objects of class Rule

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StatsCounter< Array_Type, Data_Type >	
Count stats for a single Array	117
Support < Array_Type, Data_Counter_Type, Data_Rule_Type >	
Compute the support of sufficient statistics	122
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## 4.1 File List

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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)

k genes gain 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 1081 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 711 of file phylo.hpp.

5.4 Phylo counters

#### 5.4.2.3 counter\_gains()

Functional gains for a specific function (nfun).

Definition at line 149 of file phylo.hpp.

#### 5.4.2.4 counter\_gains\_k\_offspring()

k genes gain function nfun

Definition at line 191 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 265 of file phylo.hpp.

# 5.4.2.6 counter\_longest()

Longest branch mutates (either by gain or by loss)

Definition at line 770 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 512 of file phylo.hpp.

#### 5.4.2.8 counter\_maxfuns()

Cap the number of functions per gene.

Definition at line 428 of file phylo.hpp.

#### 5.4.2.9 counter\_neofun()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 875 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 960 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 559 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 109 of file phylo.hpp.

#### 5.4.2.13 counter\_overall\_loss()

Overall functional loss.

Definition at line 382 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 625 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 32 of file typedefs.hpp.

#### 6.5.2.2 NONE

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

Definition at line 33 of file typedefs.hpp.

#### 6.5.2.3 ONE

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

Definition at line 34 of file typedefs.hpp.

# 6.5.2.4 TWO

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

Definition at line 35 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 50 of file typedefs.hpp.

#### 6.6.2.2 AS\_ZERO

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

Definition at line 49 of file typedefs.hpp.

#### 6.6.2.3 BOTH

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

Definition at line 43 of file typedefs.hpp.

#### 6.6.2.4 NONE

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

Definition at line 44 of file typedefs.hpp.

#### 6.6.2.5 ONE

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

Definition at line 45 of file typedefs.hpp.

#### 6.6.2.6 TWO

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

Definition at line 46 of file typedefs.hpp.

#### 6.6.2.7 UKNOWN

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

Definition at line 48 of file typedefs.hpp.

# **Chapter 7**

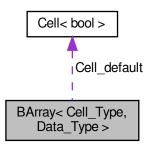
# **Class Documentation**

# 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.

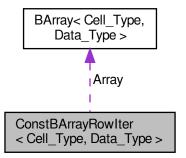
The documentation for this class was generated from the following files:

- 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\_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, std::string name_
="", std::string desc_="")

• 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 = nullptr
```

- bool delete\_data = false
- std::string name = ""
- std::string desc = ""

# 7.6.1 Detailed Description

```
\label{template} $$ \ensuremath{\sf template}$$ < 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 59 of file counters-bones.hpp.

# 7.6.2.2 Counter() [2/3]

Definition at line 61 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 76 of file counters-bones.hpp.

## 7.6.3 Member Function Documentation

## 7.6.3.1 count()

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

#### 7.6.3.2 init()

Definition at line 136 of file counters-meat.hpp.

# 7.6.3.3 operator=()

Definition at line 34 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 desc

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
std::string Counter< Array_Type, Data_Type >::desc = ""
```

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

# 7.6.4.5 init\_fun

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

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

#### 7.6.4.6 name

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
std::string Counter< Array_Type, Data_Type >::name = ""
```

Definition at line 45 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, std::string name
   —="", std::string desc\_="")
- void clear ()

# 7.7.1 Detailed Description

```
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 97 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 106 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 109 of file counters-bones.hpp.

#### 7.7.2.3 Counters() [2/2]

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

# 7.7.3 Member Function Documentation

# 7.7.3.1 add\_counter() [1/3]

Definition at line 157 of file counters-meat.hpp.

### 7.7.3.2 add\_counter() [2/3]

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

#### 7.7.3.3 add\_counter() [3/3]

Definition at line 180 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 209 of file counters-meat.hpp.

## 7.7.3.5 operator=()

Definition at line 94 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 150 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 129 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{lem:continuous} \begin{split} \text{template} &< \text{typename Cell\_Type} > \\ \text{class Entries} &< \text{Cell\_Type} > \end{split}
```

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

# **Template Parameters**

```
Cell_Type Any type
```

Definition at line 71 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 77 of file typedefs.hpp.

## 7.8.2.2 Entries() [2/2]

Definition at line 78 of file typedefs.hpp.

#### 7.8.2.3 ∼Entries()

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

Definition at line 85 of file typedefs.hpp.

# 7.8.3 Member Function Documentation

## 7.8.3.1 resize()

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

Definition at line 87 of file typedefs.hpp.

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# 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 73 of file typedefs.hpp.

## 7.8.4.2 target

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

Definition at line 74 of file typedefs.hpp.

#### 7.8.4.3 val

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

Definition at line 75 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.

Geese \* operator() (unsigned int i, bool check\_bounds=true)

Access the i-th geese element.

#### 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.

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## 7.9.2.2 ∼Flock()

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

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

# 7.9.3 Member Function Documentation

# 7.9.3.1 add\_data()

Add a tree to the flock.

#### **Parameters**

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

## 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_reduced_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.

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## 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 operator()()

```
Geese * Flock::operator() (
          unsigned int i,
          bool check_bounds = true ) [inline]
```

Access the i-th geese element.

## **Parameters**

i	Element to access	
check_bounds	When true, it will check bounds.	

Returns

 $Geese \ast$ 

Definition at line 151 of file flock-meet.hpp.

# 7.9.3.11 set\_seed()

Set the seed of the model.

# **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 ∼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()

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\_reduced\_sequence ()
- double likelihood (const std::vector< double > &par, bool as\_log=false, bool use\_reduced\_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 ()
- 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 $N$ (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

#### Geese prediction

Calculate the conditional probability

#### **Parameters**

par	Vector of parameters (terms + root).
res_prob	Vector indicating each nodes' state probability.
leave_one_out	When true, it will compute the predictions using leave-one-out, thus the prediction will be repeated nleaf times.
only_annotated	When true, it will make the predictions only on the induced sub-tree with annotated leafs.
use_reduced_sequence	Passed to the likelihood method.
preorder	For the tree traversal.

When res\_prob is specified, the function will attach the member vector probabilities from the Nodes objects. This contains the probability that the ith node has either of the possible states.

## Returns

std::vector< double > Returns the posterior probability

- std::vector< std::vector< double > > predict\_backend (const std::vector< double > &par, bool use\_
   reduced\_sequence, const std::vector< uint > &preorder)

## **Public Attributes**

- unsigned int nfunctions
- std::map< unsigned int, Node > nodes

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- barry::MapVec\_type< unsigned int > map\_to\_nodes
- std::vector< unsigned int > sequence
- std::vector< unsigned int > reduced\_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\_reduced\_sequence()

```
void Geese::calc_reduced_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.

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# 7.11.3.3 get\_probabilities()

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

Definition at line 277 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.

## 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 293 of file geese-meat.hpp.

# 7.11.3.10 nleafs()

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

Definition at line 301 of file geese-meat.hpp.

## 7.11.3.11 nnodes()

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

Definition at line 297 of file geese-meat.hpp.

# 7.11.3.12 nterms()

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

Definition at line 311 of file geese-meat.hpp.

## 7.11.3.13 observed\_counts()

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

Definition at line 318 of file geese-meat.hpp.

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#### 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()

Definition at line 165 of file geese-meat-predict.hpp.

# 7.11.3.17 predict\_backend()

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

# 7.11.3.18 print\_observed\_counts()

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

Definition at line 366 of file geese-meat.hpp.

## 7.11.3.19 set\_seed()

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

# 7.11.3.20 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.

# 7.11.3.21 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.

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# 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 map\_to\_nodes

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

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

## 7.11.4.7 nfunctions

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

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

#### 7.11.4.8 nodes

```
std::map< unsigned int, Node > Geese::nodes
```

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

### 7.11.4.9 reduced\_sequence

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

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

#### 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:

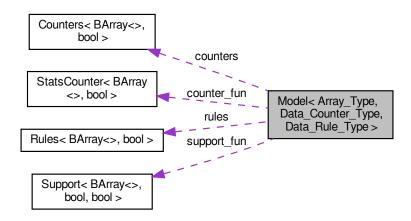
- include/barry/models/geese/geese-bones.hpp
- include/barry/models/geese/geese-meat-constructors.hpp
- include/barry/models/geese/geese-meat-likelihood.hpp
- include/barry/models/geese/geese-meat-likelihood\_exhaust.hpp
- include/barry/models/geese/geese-meat-predict.hpp
- include/barry/models/geese/geese-meat-simulate.hpp
- 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 > &params={})
- Array\_Type sample (const uint &i, const std::vector< double > &params)

## 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 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 > &params, const uint &i, bool as log=false)
- double likelihood (const std::vector< double > &params, const std::vector< double > &target\_, const uint &i, bool as log=false)
- double likelihood\_total (const std::vector< double > &params, 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.

- $\bullet \ \ \text{double } \ \ \text{get\_norm\_const} \ \ (\text{const std}:: \ \ \text{vector} < \ \ \text{double} \ > \ \& \ \ \text{params}, \ \ \text{const uint } \&i, \ \ \text{bool as\_log=false})$
- 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_\leftrightarrow Rule_Type = bool> std::vector< std::vector< Array_Type > > Model< Array_Type, Data_Counter_Type, Data_Rule_\leftrightarrow 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
- $\bullet \ \ \mathsf{std} : \! \mathsf{vector} \! < \! \mathsf{std} : \! \mathsf{vector} \! < \! \mathsf{double} > \! > \! \mathsf{vertex\_attr}$

# 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.

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#### **Parameters**

<i>vertex</i> _←	Vector of double vectors. The size equals to the number of attributes to be created. Each	l
attr_	individual vector should be of length equal to the number of vertices.	l
directed_	When true the graph as treated as directed.	l

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:



## **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

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# 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.

# 7.15.2.6 ∼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 Node Class Reference 101

## 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.

## 7.15.4.8 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

```
std::vector< bool > 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:

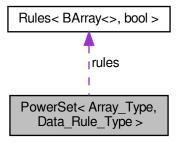
• include/barry/counters/phylo.hpp

# 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_← dat_	When true, the Rule destructor will delete the pointer, if defined.

- 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	a An object of class BArray.	
free	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:

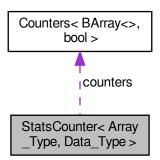
- include/barry/rules-bones.hpp
- include/barry/rules-meat.hpp

# 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

```
template<typename Array_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:

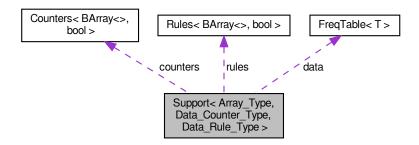
- include/barry/statscounter-bones.hpp
- include/barry/statscounter-meat.hpp

# 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**

Array←	New array over which the support will be computed.
_	

- 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
- std::vector< std::pair< uint, uint >> 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).

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#### **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.

### 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.

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### 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_\(\cho\)
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.

### 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.

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### 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 98 of file typedefs.hpp.

### 7.22.2 Member Function Documentation

### 7.22.2.1 operator()()

Definition at line 99 of file typedefs.hpp.

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

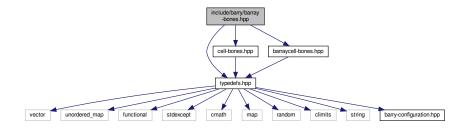
include/barry/typedefs.hpp

# **Chapter 8**

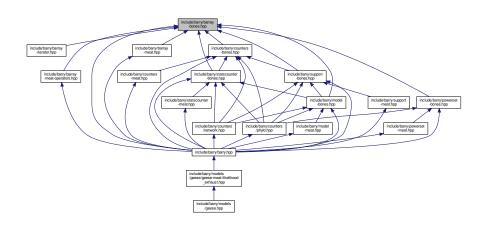
# **File Documentation**

# 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

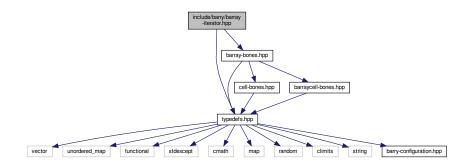
# 8.1.1.1 BARRAY\_BONES\_HPP

```
#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**

class ConstBArrayRowIter< Cell\_Type, Data\_Type >

### **Macros**

• #define BARRAY\_ITERATOR\_HPP 1

# 8.2.1 Macro Definition Documentation

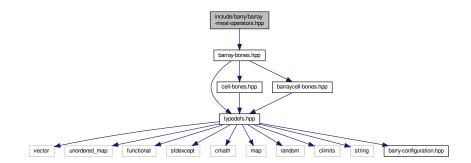
# 8.2.1.1 BARRAY\_ITERATOR\_HPP

#define BARRAY\_ITERATOR\_HPP 1

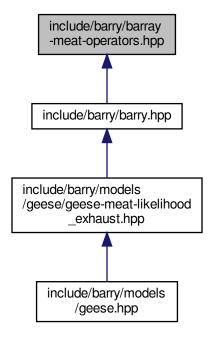
Definition at line 7 of file barray-iterator.hpp.

# 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)

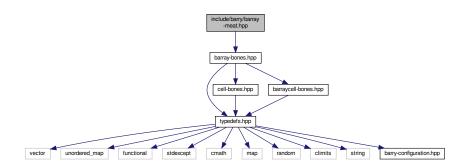
### 8.3.1 Function Documentation

# 8.3.1.1 checkdim\_()

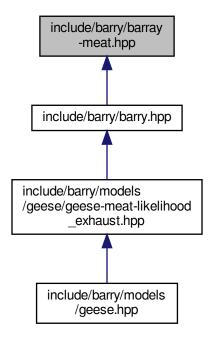
Definition at line 8 of file barray-meat-operators.hpp.

# 8.4 include/barry/barray-meat.hpp File Reference

#include "barray-bones.hpp"
Include dependency graph for barray-meat.hpp:



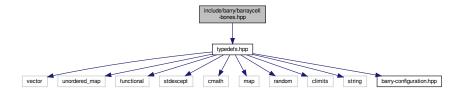
This graph shows which files directly or indirectly include this file:



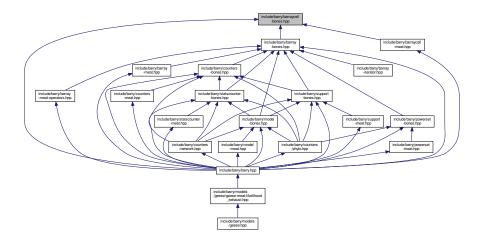
# 8.5 include/barry/barraycell-bones.hpp File Reference

#include "typedefs.hpp"

Include dependency graph for barraycell-bones.hpp:



This graph shows which files directly or indirectly include this file:

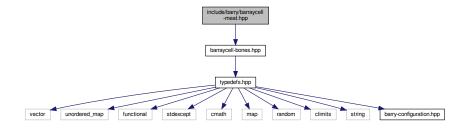


# **Classes**

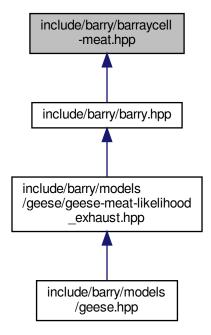
- class BArrayCell< Cell\_Type, Data\_Type >
- class BArrayCell\_const< Cell\_Type, Data\_Type >

# 8.6 include/barry/barraycell-meat.hpp File Reference

#include "barraycell-bones.hpp"
Include dependency graph for barraycell-meat.hpp:

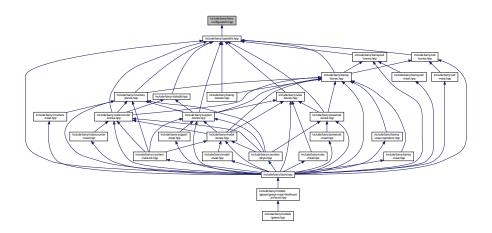


This graph shows which files directly or indirectly include this file:



# 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}/2\texttt{u})
```

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

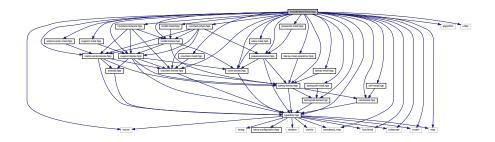
```
template<typename Ta , typename Tb >
using Map = std::map<Ta,Tb>
```

Definition at line 26 of file barry-configuration.hpp.

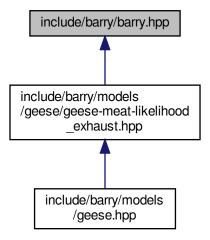
# 8.8 include/barry/barry.hpp File Reference

```
#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 )
```

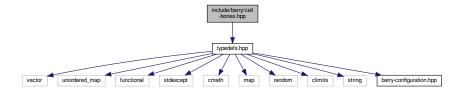
#### Value:

```
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)
```

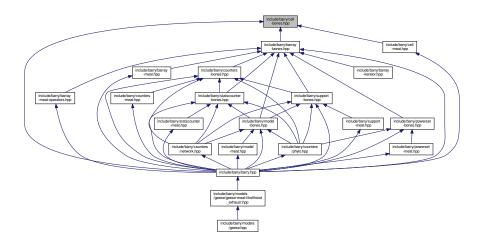
Definition at line 74 of file barry.hpp.

# 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:

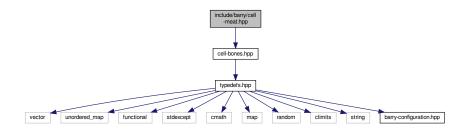


# Classes

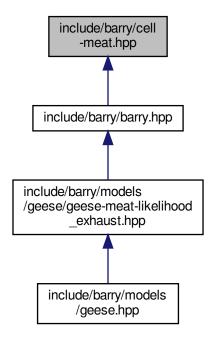
class Cell
 Cell\_Type >
 Entries in BArray. For now, it only has two members:

# 8.10 include/barry/cell-meat.hpp File Reference

#include "cell-bones.hpp"
Include dependency graph for cell-meat.hpp:



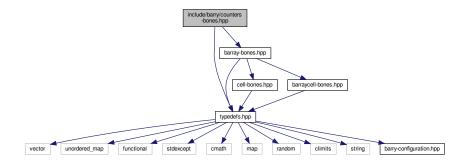
This graph shows which files directly or indirectly include this file:



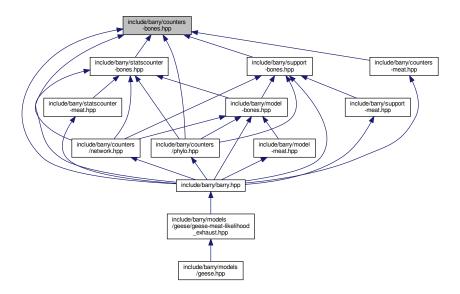
# 8.11 include/barry/col-bones.hpp File Reference

# 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:



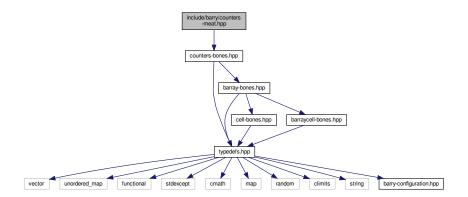
# Classes

- class Counter< Array\_Type, Data\_Type >
  - A counter function based on change statistics.
- $\bullet \ \ {\it class Counters}{< Array\_Type, \, Data\_Type} >$

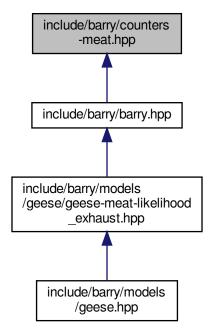
Vector of counters.

# 8.13 include/barry/counters-meat.hpp File Reference

#include "counters-bones.hpp"
Include dependency graph for counters-meat.hpp:

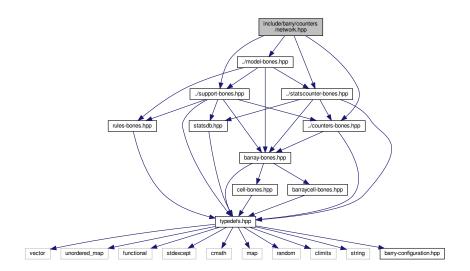


This graph shows which files directly or indirectly include this file:

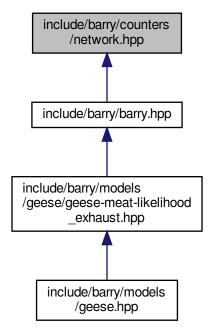


# 8.14 include/barry/counters/network.hpp File Reference

```
#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
   NetWork, NetCounterData > NetCounter
- typedef Counters < Network, NetCounterData > NetCounters
- typedef Support < Network, NetCounterData > NetSupport
- typedef StatsCounter< Network, NetCounterData > 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

### Value:

```
inline double (a) \
(const Network & Array, uint i, uint j, NetCounterData * data)
```

Function for definition of a network counter function

Definition at line 101 of file network.hpp.

# 8.14.1.4 NETWORK\_COUNTER\_LAMBDA

### Value:

```
Counter_fun_type<Network, NetCounterData> a = \
[](const Network & Array, uint i, uint j, NetCounterData * data)
```

Lambda function for definition of a network counter function

Definition at line 104 of file network.hpp.

### 8.14.1.5 NETWORK\_RULE

```
#define NETWORK_RULE(
a )
```

### Value:

```
inline bool (a) \
(const Network & Array, uint i, uint j, bool * data)
```

Function for definition of a network counter function

Definition at line 113 of file network.hpp.

### 8.14.1.6 NETWORK\_RULE\_LAMBDA

```
#define NETWORK_RULE_LAMBDA( a )
```

### Value:

```
Rule_fun_type<Network, bool> a = \
[](const Network & Array, uint i, uint j, bool * data)
```

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

```
{\tt typedef\ Counters} < {\tt Network,\ NetCounterData} > {\tt 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

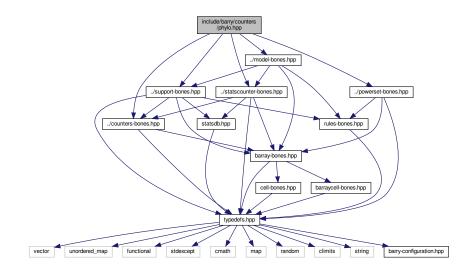
# 8.14.3.1 rules\_zerodiag()

Number of edges.

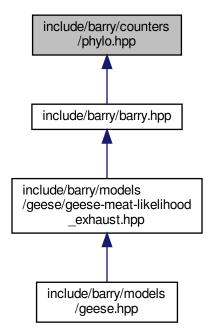
Definition at line 742 of file network.hpp.

# 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**

- std::string get\_last\_name (bool d)
- 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)

k genes gain 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.

# 8.15.3 Function Documentation

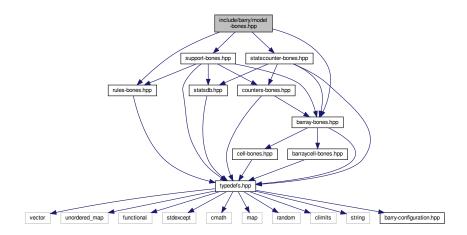
# 8.15.3.1 get\_last\_name()

```
\begin{tabular}{ll} {\tt std::string get\_last\_name (} \\ & {\tt bool } d\end{tabular} \end{tabular} \begin{tabular}{ll} {\tt inline} \end{tabular}
```

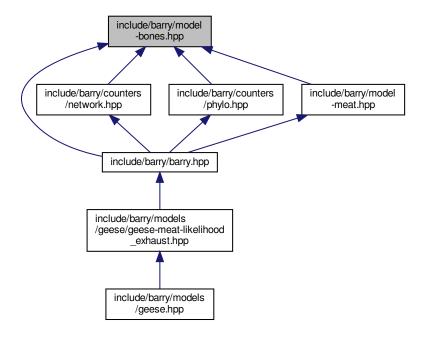
Definition at line 96 of file phylo.hpp.

# 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 > &params, const Counts\_type &support)
- double likelihood\_ (const std::vector< double > &target\_stats, const std::vector< double > &params, 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\_()

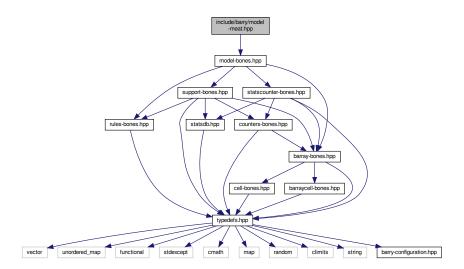
Definition at line 40 of file model-bones.hpp.

### 8.16.1.3 update\_normalizing\_constant()

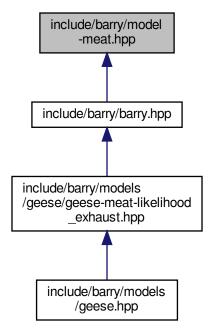
Definition at line 16 of file model-bones.hpp.

# 8.17 include/barry/model-meat.hpp File Reference

#include "model-bones.hpp"
Include dependency graph for model-meat.hpp:

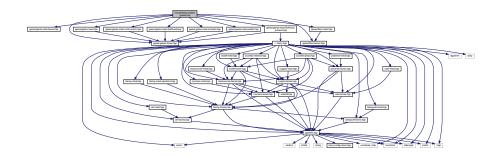


This graph shows which files directly or indirectly include this file:



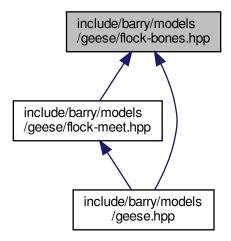
# 8.18 include/barry/models/geese.hpp File Reference

```
#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:



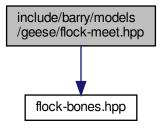
### **Classes**

· class Flock

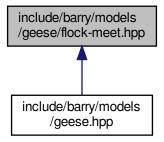
A Flock is a group of Geese.

# 8.20 include/barry/models/geese/flock-meet.hpp File Reference

#include "flock-bones.hpp"
Include dependency graph for flock-meet.hpp:

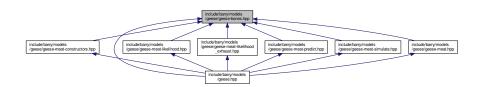


This graph shows which files directly or indirectly include this file:



# 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()

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

# 8.21.2.3 vec\_diff()

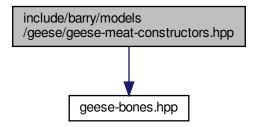
Definition at line 51 of file geese-bones.hpp.

### 8.21.2.4 vector\_caster()

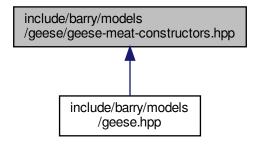
Definition at line 10 of file geese-bones.hpp.

# 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

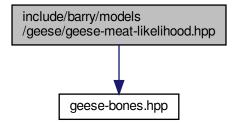
# 8.22.1.1 GEESE\_MEAT\_CONSTRUCTORS\_HPP

#define GEESE\_MEAT\_CONSTRUCTORS\_HPP 1

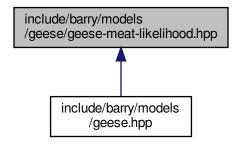
Definition at line 4 of file geese-meat-constructors.hpp.

# 8.23 include/barry/models/geese/geese-meat-likelihood.hpp File Reference

#include "geese-bones.hpp"
Include dependency graph for geese-meat-likelihood.hpp:



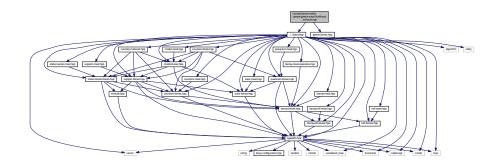
This graph shows which files directly or indirectly include this file:



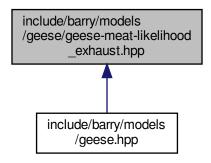
# 8.24 include/barry/models/geese/geese-meat-likelihood\_exhaust.hpp File Reference

#include "../../barry.hpp"
#include "geese-bones.hpp"

Include dependency graph for geese-meat-likelihood\_exhaust.hpp:

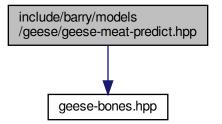


This graph shows which files directly or indirectly include this file:

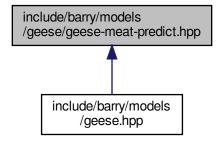


# 8.25 include/barry/models/geese/geese-meat-predict.hpp File Reference

#include "geese-bones.hpp"
Include dependency graph for geese-meat-predict.hpp:

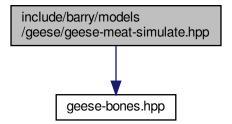


This graph shows which files directly or indirectly include this file:

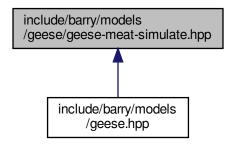


# 8.26 include/barry/models/geese/geese-meat-simulate.hpp File Reference

#include "geese-bones.hpp"
Include dependency graph for geese-meat-simulate.hpp:

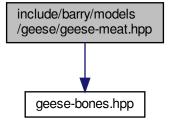


This graph shows which files directly or indirectly include this file:

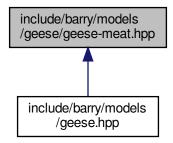


# 8.27 include/barry/models/geese/geese-meat.hpp File Reference

#include "geese-bones.hpp"
Include dependency graph for geese-meat.hpp:

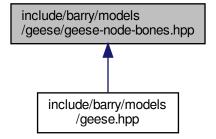


This graph shows which files directly or indirectly include this file:



# 8.28 include/barry/models/geese/geese-node-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Classes**

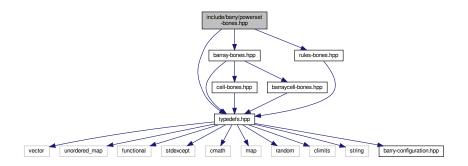
• class Node

A single node for the model.

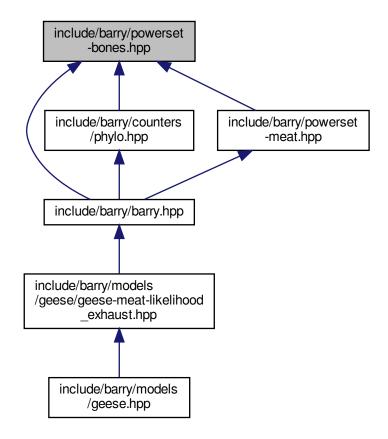
# 8.29 include/barry/powerset-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "barray-bones.hpp"
```

#include "rules-bones.hpp"
Include dependency graph for powerset-bones.hpp:



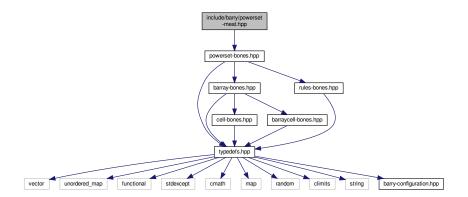
This graph shows which files directly or indirectly include this file:



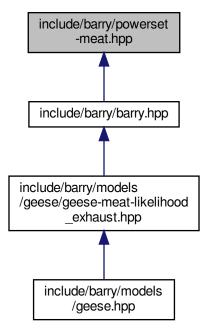
# Classes

# 8.30 include/barry/powerset-meat.hpp File Reference

#include "powerset-bones.hpp"
Include dependency graph for powerset-meat.hpp:



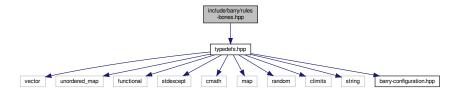
This graph shows which files directly or indirectly include this file:



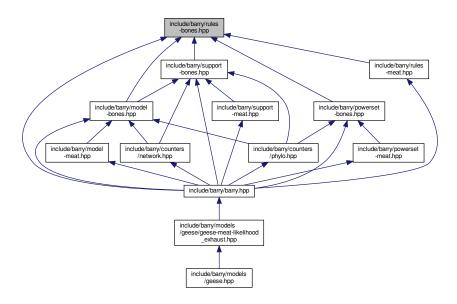
# 8.31 include/barry/rules-bones.hpp File Reference

#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.

# **Functions**

template<typename Array\_Type , typename Data\_Type >
 bool rule\_fun\_default (const Array\_Type \*array, uint i, uint j, Data\_Type \*dat)

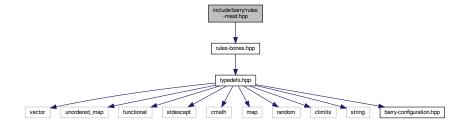
# 8.31.1 Function Documentation

#### 8.31.1.1 rule\_fun\_default()

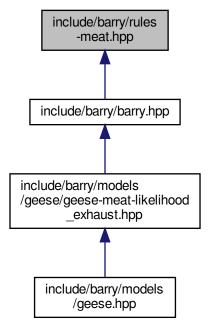
Definition at line 10 of file rules-bones.hpp.

# 8.32 include/barry/rules-meat.hpp File Reference

```
#include "rules-bones.hpp"
Include dependency graph for rules-meat.hpp:
```



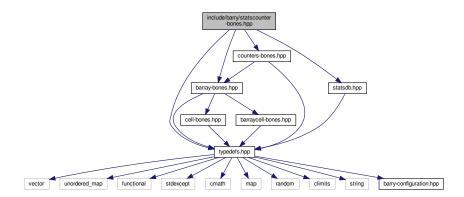
This graph shows which files directly or indirectly include this file:



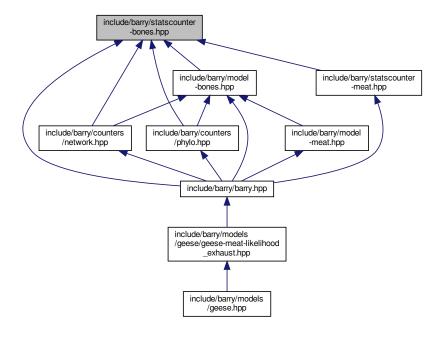
# 8.33 include/barry/statscounter-bones.hpp File Reference

```
#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:

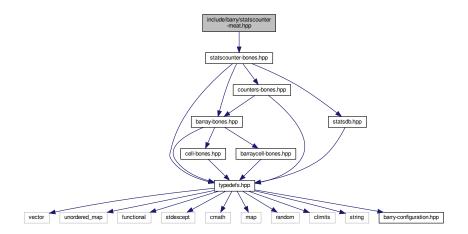


# **Classes**

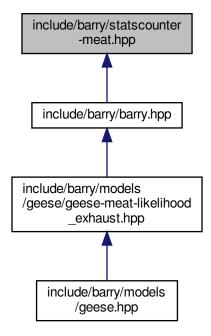
class StatsCounter < Array\_Type, Data\_Type >
 Count stats for a single Array.

# 8.34 include/barry/statscounter-meat.hpp File Reference

#include "statscounter-bones.hpp"
Include dependency graph for statscounter-meat.hpp:



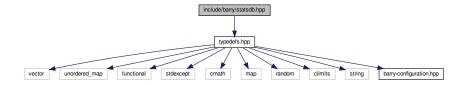
This graph shows which files directly or indirectly include this file:



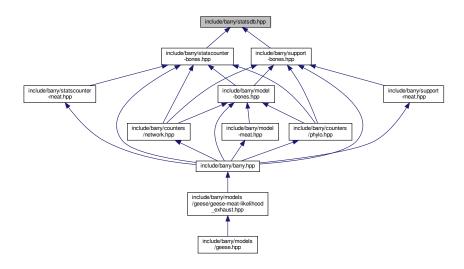
# 8.35 include/barry/statsdb.hpp File Reference

#include "typedefs.hpp"

Include dependency graph for statsdb.hpp:



This graph shows which files directly or indirectly include this file:



# **Classes**

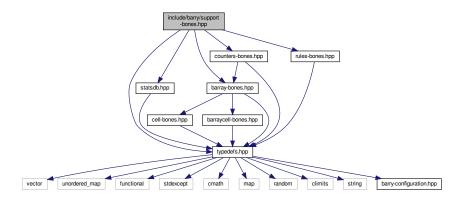
class FreqTable < T >

Database of statistics.

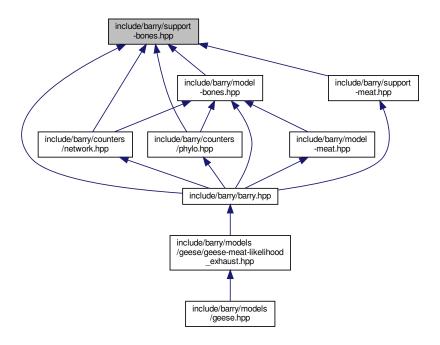
# 8.36 include/barry/support-bones.hpp File Reference

```
#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:

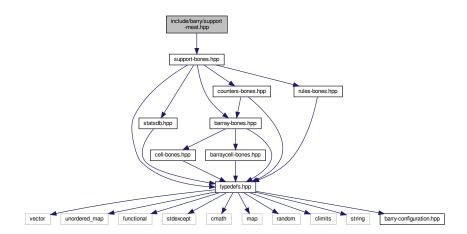


# Classes

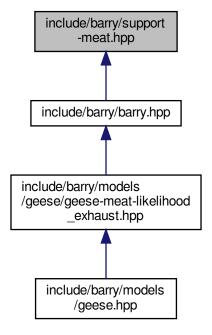
class Support < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type >
 Compute the support of sufficient statistics.

# 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

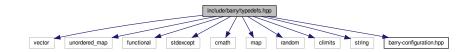
### 8.37.1.1 BARRY\_SUPPORT\_MEAT\_HPP

```
#define BARRY_SUPPORT_MEAT_HPP 1
```

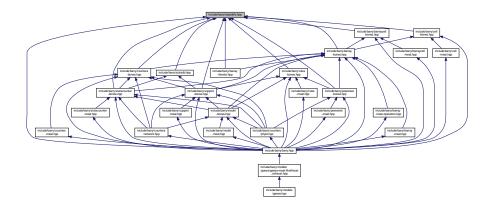
Definition at line 4 of file support-meat.hpp.

# 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 <string>
#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 124 of file typedefs.hpp.

#### 8.38.1.2 A\_ROW

```
#define A_ROW( a \ ) \ {\tt Array.el\_ij[a]}
```

Definition at line 123 of file typedefs.hpp.

### 8.38.1.3 COL

Definition at line 24 of file typedefs.hpp.

#### 8.38.1.4 ROW

```
#define ROW( a \ ) \ \  \  \text{this->el_ij[a]}
```

Definition at line 23 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 63 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 137 of file typedefs.hpp.

### 8.38.2.3 Counts\_type

```
typedef std::vector< std::pair< std::vector<double>, uint > > Counts_type
```

Definition at line 56 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 117 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 60 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 140 of file typedefs.hpp.

### 8.38.2.7 uint

```
typedef unsigned int uint
```

Definition at line 20 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 151 of file typedefs.hpp.

# 8.38.3.2 vec\_equal\_approx()

Definition at line 169 of file typedefs.hpp.

### 8.38.3.3 vec\_inner\_prod()

Definition at line 189 of file typedefs.hpp.

# 8.39 README.md File Reference

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