barry: Your go-to motif accountant 0.0-1

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## **Main Page**

### Barry: your to-go motif accountant

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

### **Examples**

#### Counting statistics in a graph

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

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

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```
netcounters::counter_ttriads(counter.counters);
netcounters::counter_isolates (counter.counters);
netcounters::counter_ctriads(counter.counters);
netcounters::counter_mutual(counter.counters);
// Counting and printing the results
std::vector< double > counts = counter.count_all();
 std::cout «
    "Edges : " « counts[0] « std::endl «
"Transitive triads : " « counts[1] « std::endl «
"Isolates : " « counts[2] « std::endl «
"C triads : " « counts[3] « std::endl «
"Mutuals : " « counts[4] « std::endl;
return 0;
```

Compiling this program using g++ g++ -std=c++11 -Wall -pedantic 08-counts.cpp -o counts && ./counts

#### Yields the following output:

```
Current view
           1
 0,] 1 1
  1,] . 1 .
 . 1
 5,]
New view
 0,] .
        1 1
[ 1,] 1 .
[ 2,] 1 .
             . 1
  3,] .
4,] 1
       . . . .
 5,] . . . .
Edges
Transitive triads : 3
       : 2
Isolates
C triads
Mutuals
```

# Namespace Index

## 2.1 Namespace List

Here is a list of all namespaces with brief descriptions:

| arry   |   |
|--|---|
| Barry: Your go-to motif accountant   | ç |
| arry::counters   |   |
| Tree class and Treelterator class  | ç |
| arry::counters::network  | ( |
| arry::counters::phylo  | ( |
| CHECK  |   |
| Integer constants used to specify which cell should be check                 | ( |
| XISTS  |   |
| Integer constants used to specify which cell should be check to exist or not | 1 |

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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   | 13 |
| BArrayCell < Cell_Type, Data_Type >  | 28 |
| BArrayCell_const< Cell_Type, Data_Type >   | 31 |
| Cell< Cell_Type >  |    |
| Entries in BArray. For now, it only has two members:   | 33 |
| ConstBArrayRowlter< Cell_Type, Data_Type >   | 38 |
| Counter< Array_Type, Data_Type >   |    |
| A counter function based on change statistics  | 40 |
| Counters< Array_Type, Data_Type >  |    |
| Vector of counters   | 43 |
| Entries < Cell_Type >  |    |
| A wrapper class to store source, target, val from a BArray object                              | 47 |
| Flock  |    |
| A Flock is a group of Geese  | 49 |
| FreqTable < T >  |    |
| Database of statistics   | 53 |
| Geese  |    |
| Annotated Phylo Model  | 55 |
| 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:                                      | 65 |
| NetCounterData   |    |
| Data class used to store arbitrary uint or double vectors                                      | 79 |
| NetworkData  |    |
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| Node   |    |
| A single node for the model  | 83 |
| NodeData   |    |
| Data definition for the PhyloArray class   | 89 |
| PowerSet < Array_Type, Data_Rule_Type >  |    |
| Powerset of a binary array   | 91 |
| Rule < Array_Type, Data_Type >   |    |
| Rule for determining if a cell should be included in a sequence                                | 97 |
| Rules < Array_Type, Data_Type >  |    |
| Vector of objects of class Rule  | 99 |

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| StatsCounter< Array_Type, Data_Type >                    |     |
|--|-----|
| Count stats for a single Array                           | 103 |
| Support< Array_Type, Data_Counter_Type, Data_Rule_Type > |     |
| Compute the support of sufficient statistics             | 108 |
| vecHasher <t></t>  | 116 |

## File Index

### 4.1 File List

Here is a list of all files with brief descriptions:

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| include/barry/barray-meat-operators.hpp                      |     |
| include/barry/barray-meat.hpp                                |     |
| include/barry/barraycell-bones.hpp                           |     |
| include/barry/barraycell-meat.hpp                            |     |
| include/barry/barry-configuration.hpp                        |     |
| include/barry/barry.hpp                                      |     |
| include/barry/cell-bones.hpp                                 |     |
| include/barry/cell-meat.hpp                                  |     |
| include/barry/col-bones.hpp                                  |     |
| include/barry/counters-bones.hpp                             |     |
| include/barry/counters-meat.hpp                              |     |
| include/barry/model-bones.hpp                                |     |
| include/barry/model-meat.hpp                                 |     |
| include/barry/powerset-bones.hpp                             |     |
| include/barry/powerset-meat.hpp                              |     |
| include/barry/rules-bones.hpp                                |     |
| include/barry/rules-meat.hpp                                 |     |
| include/barry/statscounter-bones.hpp                         |     |
| include/barry/statscounter-meat.hpp                          |     |
| include/barry/statsdb.hpp                                    |     |
| include/barry/support-bones.hpp                              |     |
| include/barry/support-meat.hpp                               |     |
| include/barry/typedefs.hpp                                   |     |
| include/barry/counters/network.hpp                           |     |
| include/barry/counters/phylo.hpp                             |     |
| include/barry/models/geese.hpp                               |     |
| include/barry/models/geese/flock-bones.hpp                   |     |
| include/barry/models/geese/flock-meet.hpp                    |     |
| 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                    |     |
| include/barry/models/geese/geese-node-bones.hpp              | 163 |
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# **Namespace Documentation**

## 5.1 barry Namespace Reference

barry: Your go-to motif accountant

#### **Namespaces**

counters

Tree class and Treelterator class.

### 5.1.1 Detailed Description

barry: Your go-to motif accountant

## 5.2 barry::counters Namespace Reference

Tree class and Treelterator class.

#### **Namespaces**

- network
- phylo

#### 5.2.1 Detailed Description

Tree class and Treelterator class.

### 5.3 barry::counters::network Namespace Reference

## 5.4 barry::counters::phylo Namespace Reference

### 5.5 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

#### 5.5.1 Detailed Description

Integer constants used to specify which cell should be check.

#### 5.5.2 Variable Documentation

#### 5.5.2.1 BOTH

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

Definition at line 33 of file typedefs.hpp.

#### 5.5.2.2 NONE

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

Definition at line 34 of file typedefs.hpp.

#### 5.5.2.3 ONE

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

Definition at line 35 of file typedefs.hpp.

### 5.5.2.4 TWO

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

Definition at line 36 of file typedefs.hpp.

### 5.6 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

#### 5.6.1 Detailed Description

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

#### 5.6.2 Variable Documentation

#### 5.6.2.1 AS\_ONE

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

Definition at line 51 of file typedefs.hpp.

#### 5.6.2.2 AS\_ZERO

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

Definition at line 50 of file typedefs.hpp.

#### 5.6.2.3 BOTH

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

Definition at line 44 of file typedefs.hpp.

#### 5.6.2.4 NONE

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

Definition at line 45 of file typedefs.hpp.

#### 5.6.2.5 ONE

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

Definition at line 46 of file typedefs.hpp.

#### 5.6.2.6 TWO

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

Definition at line 47 of file typedefs.hpp.

#### 5.6.2.7 UKNOWN

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

Definition at line 49 of file typedefs.hpp.

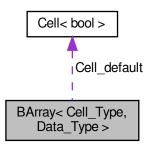
## **Class Documentation**

### 6.1 BArray < Cell\_Type, Data\_Type > Class Template Reference

Baseline class for binary arrays.

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

 $\label{lem:collaboration} \mbox{Collaboration diagram for BArray} < \mbox{Cell\_Type, Data\_Type} >:$ 



#### **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

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- 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.
- BArray (BArray < Cell\_Type, Data\_Type > &&x) noexcept
  - 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
- uint ncol () const
- uint nnozero () const

#### Cell-wise insertion/deletion

#### **Parameters**

| i,j          | Row,column  |  |
|--------------|---|--|
| check_bounds | When true and out of range, the function throws an error.     |  |
| check_exists |   |  |
|              | 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

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#### **Friends**

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

#### 6.1.1 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.

#### 6.1.2 Constructor & Destructor Documentation

#### 6.1.2.1 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.

#### 6.1.2.2 BArray() [2/6]

Empty array.

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

#### 6.1.2.3 BArray() [3/6]

Edgelist with data.

#### 6.1.2.4 BArray() [4/6]

Edgelist with no data (simpler)

#### 6.1.2.5 BArray() [5/6]

Copy constructor.

#### 6.1.2.6 BArray() [6/6]

Move operator.

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#### 6.1.2.7 ∼BArray()

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

#### 6.1.3 Member Function Documentation

#### 6.1.3.1 clear()

#### 6.1.3.2 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
```

#### 6.1.3.3 get\_cell()

#### 6.1.3.4 get\_col()

#### 6.1.3.5 get\_col\_vec() [1/2]

# 6.1.3.6 get\_col\_vec() [2/2]

#### 6.1.3.7 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>

#### 6.1.3.8 get\_row()

# 6.1.3.9 get\_row\_vec() [1/2]

#### 6.1.3.10 get\_row\_vec() [2/2]

#### 6.1.3.11 insert\_cell() [1/3]

## 6.1.3.12 insert\_cell() [2/3]

#### 6.1.3.13 insert\_cell() [3/3]

#### 6.1.3.14 is\_empty()

#### 6.1.3.15 ncol()

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

#### 6.1.3.16 nnozero()

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

## 6.1.3.17 nrow()

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

#### 6.1.3.18 operator()() [1/2]

# 6.1.3.19 operator()() [2/2]

## 6.1.3.20 operator \*=()

#### 6.1.3.21 operator+=() [1/3]

#### 6.1.3.22 operator+=() [2/3]

#### 6.1.3.23 operator+=() [3/3]

#### 6.1.3.24 operator-=() [1/3]

# 6.1.3.25 operator-=() [2/3]

#### 6.1.3.26 operator-=() [3/3]

#### 6.1.3.27 operator/=()

# 6.1.3.28 operator=() [1/2]

Move assignment.

#### 6.1.3.29 operator=() [2/2]

Assignment constructor.

## 6.1.3.30 operator==()

# 6.1.3.31 out\_of\_range()

#### 6.1.3.32 print()

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

#### 6.1.3.33 reserve()

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

# 6.1.3.34 resize()

## 6.1.3.35 rm\_cell()

# 6.1.3.36 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
```

#### 6.1.3.37 set\_data()

#### 6.1.3.38 swap\_cells()

## 6.1.3.39 swap\_cols()

#### 6.1.3.40 swap\_rows()

# 6.1.3.41 toggle\_cell()

#### 6.1.3.42 toggle\_lock()

## 6.1.3.43 transpose()

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

#### 6.1.3.44 zero col()

#### 6.1.3.45 zero\_row()

# 6.1.4 Friends And Related Function Documentation

# 6.1.4.1 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.

## 6.1.4.2 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.

# 6.1.5 Member Data Documentation

#### 6.1.5.1 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.

#### 6.1.5.2 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.

#### 6.1.5.3 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.

#### 6.1.5.4 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.

# 6.1.5.5 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.

# 6.1.5.6 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.

#### 6.1.5.7 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.

#### 6.1.5.8 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.

#### 6.1.5.9 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

# 6.2 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

# 6.2.1 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.

## 6.2.2 Constructor & Destructor Documentation

# 6.2.2.1 BArrayCell()

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

#### 6.2.2.2 ∼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.

# 6.2.3 Member Function Documentation

#### 6.2.3.1 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.

#### 6.2.3.2 operator\*=()

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

# 6.2.3.3 operator+=()

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

## 6.2.3.4 operator-=()

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

## 6.2.3.5 operator/=()

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

# 6.2.3.6 operator=()

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

#### 6.2.3.7 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

# 6.3 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 j\_, bool check\_bounds=true)
- ∼BArrayCell const ()
- operator Cell\_Type () const
- bool operator== (const Cell\_Type &val) const
- bool operator!= (const Cell\_Type &val) const
- $\bullet \ \ bool \ \ \ operator < (const \ Cell\_Type \ \&val) \ const$
- bool operator> (const Cell\_Type &val) const
- bool operator<= (const Cell\_Type &val) const
- bool operator>= (const Cell\_Type &val) const

# 6.3.1 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.

#### 6.3.2 Constructor & Destructor Documentation

# 6.3.2.1 BArrayCell\_const()

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

#### 6.3.2.2 ∼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.

#### 6.3.3 Member Function Documentation

## 6.3.3.1 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.

## 6.3.3.2 operator"!=()

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

#### 6.3.3.3 operator<()

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

#### 6.3.3.4 operator<=()

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

#### 6.3.3.5 operator==()

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

#### 6.3.3.6 operator>()

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

# 6.3.3.7 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

# 6.4 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

# 6.4.1 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.

#### 6.4.2 Constructor & Destructor Documentation

#### 6.4.2.1 Cell() [1/7]

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

# 6.4.2.2 Cell() [2/7]

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

# 6.4.2.3 ∼CeII()

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

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

# 6.4.2.4 Cell() [3/7]

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

# 6.4.2.5 Cell() [4/7]

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

# 6.4.2.6 Cell() [5/7]

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

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

# 6.4.2.7 Cell() [6/7]

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

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

## 6.4.2.8 Cell() [7/7]

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

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

## 6.4.3 Member Function Documentation

## 6.4.3.1 add() [1/4]

## 6.4.3.2 add() [2/4]

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

# 6.4.3.3 add() [3/4]

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

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

## 6.4.3.4 add() [4/4]

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

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

# 6.4.3.5 operator Cell\_Type()

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

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

## 6.4.3.6 operator=() [1/2]

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

# 6.4.3.7 operator=() [2/2]

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

# 6.4.4 Member Data Documentation

# 6.4.4.1 value

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

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

## 6.4.4.2 visited

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

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

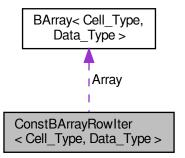
The documentation for this class was generated from the following files:

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

# 6.5 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\_)
- ∼ConstBArrayRowIter ()

# **Public Attributes**

- uint current\_row
- uint current\_col
- Row\_type< Cell\_Type >::const\_iterator iter
- const BArray< Cell\_Type, Data\_Type > \* Array

# 6.5.1 Detailed Description

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

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

# 6.5.2 Constructor & Destructor Documentation

# 6.5.2.1 ConstBArrayRowIter()

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

#### 6.5.2.2 ∼ConstBArrayRowlter()

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

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

#### 6.5.3 Member Data Documentation

#### 6.5.3.1 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.

## 6.5.3.2 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.

#### 6.5.3.3 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.

#### 6.5.3.4 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

# 6.6 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_←<br>data_ | When true, the destructor will delete the pointer in the main data.  |

- Counter ()
- Counter (Counter\_fun\_type< Array\_Type, Data\_Type > count\_fun\_, Counter\_fun\_type< Array\_Type,
   Data\_Type > init\_fun\_=nullptr, Data\_Type \*data\_=nullptr, bool delete\_data\_=false)

Counter (const Counter< Array\_Type, Data\_Type > &counter\_)

## **Public Attributes**

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

- Data Type \* data = nullptr
- bool delete\_data = false

# 6.6.1 Detailed Description

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

A counter function based on change statistics.

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

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

## 6.6.2 Constructor & Destructor Documentation

# 6.6.2.1 Counter() [1/3]

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

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

# 6.6.2.2 Counter() [2/3]

Definition at line 36 of file counters-bones.hpp.

#### 6.6.2.3 Counter() [3/3]

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

#### 6.6.2.4 ~Counter()

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

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

## 6.6.3 Member Function Documentation

#### 6.6.3.1 count()

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

#### 6.6.3.2 init()

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

# 6.6.3.3 operator=()

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

#### 6.6.4 Member Data Documentation

## 6.6.4.1 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 18 of file counters-bones.hpp.

#### 6.6.4.2 data

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

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

#### 6.6.4.3 delete\_data

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

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

#### 6.6.4.4 init fun

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

Definition at line 19 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

# 6.7 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.

• uint size () const

Number of counters in the set.

- void add\_counter (Counter< Array\_Type, Data\_Type > &counter)
- void add\_counter (Counter< Array\_Type, Data\_Type > \*counter)
- void add\_counter (Counter\_fun\_type< Array\_Type, Data\_Type > count\_fun\_, Counter\_fun\_type< Array\_
   —
   Type, Data\_Type > init\_fun\_=nullptr, Data\_Type \*data\_=nullptr, bool delete\_data\_=false)
- void clear ()

# 6.7.1 Detailed Description

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

Vector of counters.

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

Definition at line 69 of file counters-bones.hpp.

#### 6.7.2 Constructor & Destructor Documentation

#### 6.7.2.1 Counters() [1/2]

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

Definition at line 78 of file counters-bones.hpp.

#### 6.7.2.2 ∼Counters()

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

Definition at line 81 of file counters-bones.hpp.

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

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

#### 6.7.3 Member Function Documentation

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

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

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

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

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

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

#### 6.7.3.4 clear()

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

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

#### 6.7.3.5 operator=()

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

# 6.7.3.6 operator[]()

Returns a pointer to a particular counter.

#### **Parameters**

```
idx Id of the counter
```

#### Returns

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

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

# 6.7.3.7 size()

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

Number of counters in the set.

Returns

uint

Definition at line 101 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

# 6.8 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

# 6.8.1 Detailed Description

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

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

**Template Parameters** 

```
Cell_Type Any type
```

Definition at line 72 of file typedefs.hpp.

## 6.8.2 Constructor & Destructor Documentation

#### 6.8.2.1 Entries() [1/2]

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

Definition at line 78 of file typedefs.hpp.

#### 6.8.2.2 Entries() [2/2]

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

Definition at line 79 of file typedefs.hpp.

# 6.8.2.3 $\sim$ Entries()

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

Definition at line 86 of file typedefs.hpp.

## 6.8.3 Member Function Documentation

## 6.8.3.1 resize()

Definition at line 88 of file typedefs.hpp.

# 6.8.4 Member Data Documentation

#### 6.8.4.1 source

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

Definition at line 74 of file typedefs.hpp.

# 6.8.4.2 target

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

Definition at line 75 of file typedefs.hpp.

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#### 6.8.4.3 val

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

Definition at line 76 of file typedefs.hpp.

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

• include/barry/typedefs.hpp

# 6.9 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)
- void set seed (const unsigned int &s)
- void init ()
- phylocounters::PhyloCounters \* counters\_ptr ()

#### Information about the model

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

# 6.9.1 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.

Definition at line 12 of file flock-bones.hpp.

# 6.9.2 Constructor & Destructor Documentation

## 6.9.2.1 Flock()

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

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

#### 6.9.2.2 ∼Flock()

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

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

## 6.9.3 Member Function Documentation

# 6.9.3.1 add\_data()

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

# 6.9.3.2 counters\_ptr()

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

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

# 6.9.3.3 init()

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

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

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# 6.9.3.4 likelihood\_joint()

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

# 6.9.3.5 nfuns()

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

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

# 6.9.3.6 nleafs()

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

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

# 6.9.3.7 nnodes()

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

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

#### 6.9.3.8 nterms()

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

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

## 6.9.3.9 ntrees()

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

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

# 6.9.3.10 set\_seed()

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

# 6.9.4 Member Data Documentation

## 6.9.4.1 dat

```
std::vector< Geese > Flock::dat
```

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

#### 6.9.4.2 initialized

```
bool Flock::initialized = false
```

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

# 6.9.4.3 nfunctions

```
unsigned int Flock::nfunctions = Ou
```

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

#### 6.9.4.4 rengine

std::mt19937 Flock::rengine

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

#### 6.9.4.5 support

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

Definition at line 21 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

# **6.10** 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

# 6.10.1 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.

## 6.10.2 Constructor & Destructor Documentation

# 6.10.2.1 FreqTable()

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

Definition at line 28 of file statsdb.hpp.

# 6.10.2.2 $\sim$ FreqTable()

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

Definition at line 29 of file statsdb.hpp.

# 6.10.3 Member Function Documentation

# 6.10.3.1 add()

Definition at line 46 of file statsdb.hpp.

# 6.10.3.2 as\_vector()

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

Definition at line 60 of file statsdb.hpp.

# 6.10.3.3 clear()

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

Definition at line 82 of file statsdb.hpp.

# 6.10.3.4 get\_data()

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

Definition at line 72 of file statsdb.hpp.

# 6.10.3.5 get\_data\_ptr()

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

Definition at line 77 of file statsdb.hpp.

# 6.10.3.6 print()

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

Definition at line 101 of file statsdb.hpp.

# 6.10.3.7 reserve()

```
template<typename T >
void FreqTable< T >::reserve (
         unsigned int n ) [inline]
```

Definition at line 88 of file statsdb.hpp.

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

• include/barry/statsdb.hpp

# 6.11 Geese Class Reference

Annotated Phylo Model.

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

#### **Public Member Functions**

- ∼Geese ()
- · void init ()
- void inherit support (const Geese &model, bool delete support =false)
- void calc\_sequence (Node \*n=nullptr)
- void calc likelihood sequence ()
- double likelihood (const std::vector< double > &par, bool as\_log=false, bool use\_likelihood\_sequence=true)
- double likelihood\_exhaust (const std::vector< double > &par)
- std::vector< double > get\_probabilities () const
- void set\_seed (const unsigned int &s)
- std::vector< std::vector< unsigned int > > simulate (const std::vector< double > &par)
- std::vector< std::vector< double >> observed\_counts ()
- void print\_observed\_counts ()
- std::vector< std::vector< double > > predict (const std::vector< double > &p, std::vector< std::vector< double > > \*res\_prob=nullptr)

Calculate the conditional probability.

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

## Construct a new Geese object

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

#### **Parameters**

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

- 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
- unsigned int nnodes () const
- unsigned int nleafs () const
- unsigned int nterms () const

## **Public Attributes**

- unsigned int nfunctions
- barry::Map< unsigned int, Node > nodes
- barry::MapVec type< unsigned int > map to nodes
- std::vector< unsigned int > sequence
- std::vector< unsigned int > likelihood\_sequence

- bool initialized = false
- bool delete\_rengine = false
- bool delete\_counters = false
- bool delete\_support = false

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

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

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

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

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

# 6.11.1 Detailed Description

Annotated Phylo Model.

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

# 6.11.2 Constructor & Destructor Documentation

# 6.11.2.1 Geese() [1/4]

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

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

# 6.11.2.2 Geese() [2/4]

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

# 6.11.2.3 Geese() [3/4]

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

# 6.11.2.4 Geese() [4/4]

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

## 6.11.2.5 ∼Geese()

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

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

# 6.11.3 Member Function Documentation

# 6.11.3.1 calc\_likelihood\_sequence()

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

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

# 6.11.3.2 calc\_sequence()

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

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

# 6.11.3.3 get\_probabilities()

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

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

#### 6.11.3.4 inherit\_support()

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

# 6.11.3.5 init()

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

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

# 6.11.3.6 init\_node()

```
void Geese::init_node (
          Node & n ) [inline]
```

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

# 6.11.3.7 likelihood()

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

# 6.11.3.8 likelihood\_exhaust()

Definition at line 7 of file geese-meat-likelihood\_exhaust.hpp.

# 6.11.3.9 nfuns()

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

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

# 6.11.3.10 nleafs()

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

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

# 6.11.3.11 nnodes()

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

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

# 6.11.3.12 nterms()

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

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

# 6.11.3.13 observed\_counts()

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

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

# 6.11.3.14 operator=() [1/2]

# 6.11.3.15 operator=() [2/2]

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

# 6.11.3.16 predict()

Calculate the conditional probability.

#### **Parameters**

```
p Vector of parameters
```

# Returns

std::vector< double > Returns the posterior probability

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

#### 6.11.3.17 print\_observed\_counts()

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

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

# 6.11.3.18 set\_seed()

```
void Geese::set_seed ( {\tt const\ unsigned\ int\ \&\ s\ )} \quad [{\tt inline}]
```

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

# 6.11.3.19 simulate()

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

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

# 6.11.3.20 update\_annotations()

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

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

# 6.11.4 Member Data Documentation

#### 6.11.4.1 counters

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

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

# 6.11.4.2 delete\_counters

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

Definition at line 99 of file geese-bones.hpp.

# 6.11.4.3 delete\_rengine

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

Definition at line 98 of file geese-bones.hpp.

# 6.11.4.4 delete\_support

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

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

#### 6.11.4.5 initialized

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

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

# 6.11.4.6 likelihood\_sequence

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

Definition at line 94 of file geese-bones.hpp.

# 6.11.4.7 map\_to\_nodes

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

Definition at line 90 of file geese-bones.hpp.

## 6.11.4.8 nfunctions

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

Definition at line 88 of file geese-bones.hpp.

#### 6.11.4.9 nodes

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

Definition at line 89 of file geese-bones.hpp.

# 6.11.4.10 rengine

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

Definition at line 81 of file geese-bones.hpp.

#### 6.11.4.11 sequence

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

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

#### 6.11.4.12 states

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

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

# 6.11.4.13 support

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

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

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

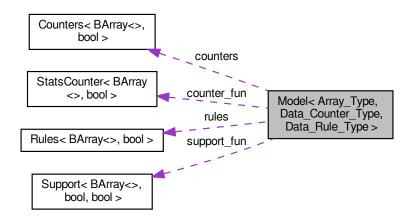
- include/barry/models/geese/geese-bones.hpp
- include/barry/models/geese/geese-meat-constructors.hpp
- include/barry/models/geese/geese-meat-likelihood.hpp
- include/barry/models/geese/geese-meat-likelihood\_exhaust.hpp
- include/barry/models/geese/geese-meat-predict.hpp
- include/barry/models/geese/geese-meat-simulate.hpp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

# 6.12 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 ()
- 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 add\_counter (Counter\_fun\_type < Array\_Type, Data\_Counter\_Type > count\_fun\_, Counter\_fun\_type <
   Array\_Type, Data\_Counter\_Type > init\_fun\_=nullptr, Data\_Counter\_Type \*data\_=nullptr, bool delete\_
   data\_=false)

void set\_counters (Counters < Array\_Type, Data\_Counter\_Type > \*counters\_)

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

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

- void add\_rule (Rule < Array\_Type, Data\_Rule\_Type > &rule)
- void add rule (Rule < Array Type, Data Rule Type > \*rule)
- void add\_rule (Rule\_fun\_type< Array\_Type, Data\_Rule\_Type > count\_fun\_, Data\_Rule\_Type \*data\_←
   =nullptr, bool delete data =false)
- void set\_rules (Rules < Array\_Type, Data\_Rule\_Type > \*rules\_)

#### Likelihood functions.

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

#### **Parameters**

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

- double likelihood (const std::vector< double > &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.     |

- double get\_norm\_const (const std::vector< double > &params, const uint &i, bool as\_log=false)
- const std::vector< Array\_Type > \* get\_pset (const uint &i)
- const std::vector< std::vector< double > > \* get\_stats (const uint &i)

## Size of the model

Number of different supports included in the model

This will return the size of stats.

#### Returns

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

- unsigned int size () const
- unsigned int size\_unique () const
- unsigned int nterms () const

#### **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)

# 6.12.1 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 95 of file model-bones.hpp.

# 6.12.2 Constructor & Destructor Documentation

# 6.12.2.1 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.

# 6.12.2.2 Model() [2/3]

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

#### 6.12.2.3 Model() [3/3]

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

# 6.12.2.4 ∼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 190 of file model-bones.hpp.

# 6.12.3 Member Function Documentation

#### 6.12.3.1 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.

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

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

#### 6.12.3.3 add counter() [2/3]

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

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

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

# 6.12.3.5 add\_rule() [1/3]

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

#### 6.12.3.6 add\_rule() [2/3]

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

#### 6.12.3.7 add\_rule() [3/3]

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

# 6.12.3.8 get\_norm\_const()

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

# 6.12.3.9 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.

# 6.12.3.10 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.

# 6.12.3.11 likelihood() [1/3]

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

#### 6.12.3.12 likelihood() [2/3]

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

#### 6.12.3.13 likelihood() [3/3]

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

# 6.12.3.14 likelihood\_total()

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

#### 6.12.3.15 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]
```

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

## 6.12.3.16 operator=()

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

#### 6.12.3.17 print\_stats()

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

# 6.12.3.18 sample() [1/2]

#### 6.12.3.19 sample() [2/2]

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

## 6.12.3.20 set\_counters()

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

#### 6.12.3.21 set keygen()

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

# 6.12.3.22 set\_rengine()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool>
void Model< Array_Type, Data_Counter_Type, Data_Rule_Type >::set_rengine (
    std::mt19937 * rengine_,
    bool delete_ = false ) [inline]
```

Definition at line 106 of file model-bones.hpp.

# 6.12.3.23 set\_rules()

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

#### 6.12.3.24 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 116 of file model-bones.hpp.

## 6.12.3.25 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]
```

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

#### 6.12.3.26 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]
```

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

# 6.12.3.27 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]
```

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

## 6.12.4 Member Data Documentation

#### 6.12.4.1 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 154 of file model-bones.hpp.

#### 6.12.4.2 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 155 of file model-bones.hpp.

## 6.12.4.3 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 172 of file model-bones.hpp.

#### 6.12.4.4 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 169 of file model-bones.hpp.

#### 6.12.4.5 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 105 of file model-bones.hpp.

# 6.12.4.6 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 178 of file model-bones.hpp.

# 6.12.4.7 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 182 of file model-bones.hpp.

## 6.12.4.8 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 162 of file model-bones.hpp.

#### 6.12.4.9 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 131 of file model-bones.hpp.

## 6.12.4.10 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 177 of file model-bones.hpp.

# 6.12.4.11 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 176 of file model-bones.hpp.

#### 6.12.4.12 pset\_arrays

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

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

# 6.12.4.13 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 142 of file model-bones.hpp.

#### 6.12.4.14 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 141 of file model-bones.hpp.

# 6.12.4.15 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 104 of file model-bones.hpp.

# 6.12.4.16 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 170 of file model-bones.hpp.

#### 6.12.4.17 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 130 of file model-bones.hpp.

# 6.12.4.18 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 171 of file model-bones.hpp.

# 6.12.4.19 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 153 of file model-bones.hpp.

# 6.12.4.20 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 139 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

# 6.13 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

# 6.13.1 Detailed Description

Data class used to store arbitrary uint or double vectors.

Definition at line 50 of file network.hpp.

#### 6.13.2 Constructor & Destructor Documentation

# 6.13.2.1 NetCounterData() [1/2]

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

Definition at line 56 of file network.hpp.

# 6.13.2.2 NetCounterData() [2/2]

Definition at line 57 of file network.hpp.

# 6.13.2.3 ∼NetCounterData()

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

Definition at line 62 of file network.hpp.

# 6.13.3 Member Data Documentation

#### 6.13.3.1 indices

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

Definition at line 53 of file network.hpp.

# 6.13.3.2 numbers

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

Definition at line 54 of file network.hpp.

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

• include/barry/counters/network.hpp

# 6.14 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
- $\bullet \ \ \mathsf{std} : \! \mathsf{vector} \! < \! \mathsf{std} : \! \mathsf{vector} \! < \! \mathsf{double} > \! > \! \mathsf{vertex\_attr}$

# 6.14.1 Detailed Description

Data class for Networks.

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

Definition at line 15 of file network.hpp.

# 6.14.2 Constructor & Destructor Documentation

# 6.14.2.1 NetworkData() [1/3]

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

Definition at line 21 of file network.hpp.

# 6.14.2.2 NetworkData() [2/3]

Constructor using a single attribute.

#### **Parameters**

| vertex_←<br>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 28 of file network.hpp.

# 6.14.2.3 NetworkData() [3/3]

```
NetworkData::NetworkData (
          std::vector< std::vector< double > > vertex_attr_,
          bool directed_ = true ) [inline]
```

Constructor using multiple attributes.

6.15 Node Class Reference 83

#### **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 39 of file network.hpp.

# 6.14.2.4 ~NetworkData()

NetworkData::~NetworkData ( ) [inline]

Definition at line 45 of file network.hpp.

# 6.14.3 Member Data Documentation

#### 6.14.3.1 directed

bool NetworkData::directed = true

Definition at line 18 of file network.hpp.

# 6.14.3.2 vertex\_attr

std::vector< std::vector< double > > NetworkData::vertex\_attr

Definition at line 19 of file network.hpp.

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

• include/barry/counters/network.hpp

# 6.15 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

#### 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.

# 6.15.1 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.

## 6.15.2 Constructor & Destructor Documentation

6.15 Node Class Reference 85

# 6.15.2.1 Node() [1/5]

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

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

# 6.15.2.2 Node() [2/5]

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

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

# 6.15.2.3 Node() [3/5]

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

# 6.15.2.4 Node() [4/5]

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

# **6.15.2.5** Node() [5/5]

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

# 6.15.2.6 ∼Node()

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

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

# 6.15.3 Member Function Documentation

# 6.15.3.1 get\_parent()

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

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

# 6.15.3.2 is\_leaf()

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

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

# 6.15.4 Member Data Documentation

# 6.15.4.1 annotations

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

Observed annotations (only defined for Geese)

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

# 6.15.4.2 array

phylocounters::PhyloArray Node::array

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

6.15 Node Class Reference 87

# 6.15.4.3 arrays

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

Arrays given all possible states.

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

# 6.15.4.4 duplication

```
bool Node::duplication
```

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

# 6.15.4.5 id

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

Id of the node (as specified in the input)

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

# 6.15.4.6 narray

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

ID of the array in the model.

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

# 6.15.4.7 offspring

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

Offspring nodes.

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

# 6.15.4.8 ord

```
unsigned int Node::ord
```

Order in which the node was created.

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

# 6.15.4.9 parent

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

Parent node.

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

# 6.15.4.10 probability

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

The probability of observing each state.

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

#### 6.15.4.11 subtree prob

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

Induced subtree probabilities.

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

#### 6.15.4.12 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

# 6.16 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

# 6.16.1 Detailed Description

Data definition for the PhyloArray class.

This holds basic information about a given node.

Definition at line 16 of file phylo.hpp.

# 6.16.2 Constructor & Destructor Documentation

#### 6.16.2.1 NodeData() [1/2]

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

Definition at line 34 of file phylo.hpp.

# 6.16.2.2 NodeData() [2/2]

Definition at line 36 of file phylo.hpp.

# 6.16.2.3 ∼NodeData()

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

Definition at line 42 of file phylo.hpp.

# 6.16.3 Member Data Documentation

# 6.16.3.1 blengths

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

Branch length.

Definition at line 22 of file phylo.hpp.

# 6.16.3.2 duplication

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

Definition at line 32 of file phylo.hpp.

## 6.16.3.3 states

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

State of the parent node.

Definition at line 27 of file phylo.hpp.

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

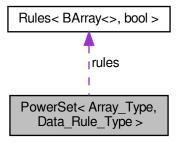
• include/barry/counters/phylo.hpp

# 6.17 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 ()
- uint size () const
- 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

#### 6.17.1 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.

#### 6.17.2 Constructor & Destructor Documentation

#### 6.17.2.1 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.

#### 6.17.2.2 PowerSet() [2/3]

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

#### 6.17.2.3 PowerSet() [3/3]

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

#### 6.17.2.4 ∼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.

#### 6.17.3 Member Function Documentation

#### 6.17.3.1 add rule() [1/3]

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

#### 6.17.3.2 add rule() [2/3]

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

#### 6.17.3.3 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.

#### 6.17.3.4 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.

#### 6.17.3.5 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.

#### 6.17.3.6 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.

#### 6.17.3.7 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.

#### 6.17.3.8 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.

#### 6.17.3.9 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.

#### 6.17.3.10 operator[]()

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

#### 6.17.3.11 reset()

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

#### 6.17.3.12 size()

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

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

#### 6.17.4 Member Data Documentation

#### 6.17.4.1 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.

#### 6.17.4.2 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.

#### 6.17.4.3 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.

#### 6.17.4.4 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.

#### 6.17.4.5 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.

#### 6.17.4.6 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.

#### 6.17.4.7 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.

#### 6.17.4.8 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

## 6.18 Rule < Array\_Type, Data\_Type > Class Template Reference

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

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

#### **Public Member Functions**

- ∼Rule ()
- bool locked (const Array\_Type &a, uint i, uint j)

#### Construct a new Rule object

Construct a new Rule object

#### **Parameters**

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

- Rule ()
- Rule (Rule\_fun\_type< Array\_Type, Data\_Type > fun\_, Data\_Type \*dat\_=nullptr, bool delete\_dat\_=false)

#### 6.18.1 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.

#### 6.18.2 Constructor & Destructor Documentation

#### 6.18.2.1 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.

#### 6.18.2.2 Rule() [2/2]

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

#### 6.18.2.3 ∼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.

#### 6.18.3 Member Function Documentation

#### 6.18.3.1 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

## 6.19 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
- 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)

#### 6.19.1 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.

#### 6.19.2 Constructor & Destructor Documentation

#### 6.19.2.1 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.

#### 6.19.2.2 Rules() [2/2]

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

#### 6.19.2.3 ∼Rules()

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

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

#### 6.19.3 Member Function Documentation

#### 6.19.3.1 add\_rule() [1/3]

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

#### 6.19.3.2 add\_rule() [2/3]

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

#### 6.19.3.3 add\_rule() [3/3]

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

#### 6.19.3.4 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.

#### 6.19.3.5 get\_seq()

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

#### **Parameters**

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

#### Returns

Nothing.

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

#### 6.19.3.6 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.

#### 6.19.3.7 operator=()

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

#### 6.19.3.8 size()

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

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

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

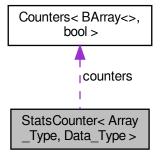
- include/barry/rules-bones.hpp
- include/barry/rules-meat.hpp

## 6.20 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
- Counters < Array\_Type, Data\_Type > \* counters
- bool counter\_deleted = false

#### 6.20.1 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.

#### 6.20.2 Constructor & Destructor Documentation

#### 6.20.2.1 StatsCounter() [1/2]

Creator of a StatsCounter

#### **Parameters**

| Array← | A const pointer to a BArray. |
|--------|------------------------------|
|        |                              |

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

#### 6.20.2.2 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.

#### 6.20.2.3 ∼StatsCounter()

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

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

#### 6.20.3 Member Function Documentation

#### 6.20.3.1 add\_counter() [1/2]

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

#### 6.20.3.2 add counter() [2/2]

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

#### 6.20.3.3 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.

#### 6.20.3.4 count\_current()

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

#### 6.20.3.5 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.

#### 6.20.3.6 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.

#### 6.20.3.7 set\_counters()

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

#### 6.20.4 Member Data Documentation

#### 6.20.4.1 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.

#### 6.20.4.2 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.

#### 6.20.4.3 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.

#### 6.20.4.4 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.

#### 6.20.4.5 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:

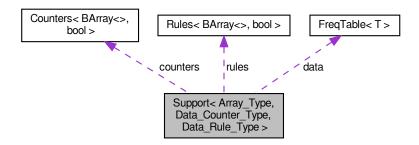
- include/barry/statscounter-bones.hpp
- include/barry/statscounter-meat.hpp

# 6.21 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)

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
- 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

#### 6.21.1 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.

#### 6.21.2 Constructor & Destructor Documentation

#### 6.21.2.1 Support() [1/3]

Constructor passing a reference Array.

Definition at line 55 of file support-bones.hpp.

#### 6.21.2.2 Support() [2/3]

Constructor specifying the dimensions of the array (empty).

Definition at line 63 of file support-bones.hpp.

#### 6.21.2.3 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 69 of file support-bones.hpp.

#### 6.21.2.4 ∼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 75 of file support-bones.hpp.

#### 6.21.3 Member Function Documentation

#### 6.21.3.1 add\_counter() [1/2]

Definition at line 171 of file support-meat.hpp.

#### 6.21.3.2 add\_counter() [2/2]

Definition at line 181 of file support-meat.hpp.

#### 6.21.3.3 add\_rule() [1/2]

Definition at line 208 of file support-meat.hpp.

#### 6.21.3.4 add\_rule() [2/2]

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

#### 6.21.3.5 calc()

Computes the entire support.

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

#### **Parameters**

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

Definition at line 153 of file support-meat.hpp.

#### 6.21.3.6 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 245 of file support-meat.hpp.

#### 6.21.3.7 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 252 of file support-meat.hpp.

#### 6.21.3.8 init\_support()

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

#### 6.21.3.9 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 259 of file support-meat.hpp.

#### 6.21.3.10 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 69 of file support-meat.hpp.

#### 6.21.3.11 reset\_array() [2/2]

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

#### 6.21.3.12 set counters()

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

#### 6.21.3.13 set\_rules()

Definition at line 228 of file support-meat.hpp.

#### 6.21.4 Member Data Documentation

#### 6.21.4.1 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 51 of file support-bones.hpp.

#### 6.21.4.2 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 49 of file support-bones.hpp.

#### 6.21.4.3 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 50 of file support-bones.hpp.

#### 6.21.4.4 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.

#### 6.21.4.5 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.

#### 6.21.4.6 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 48 of file support-bones.hpp.

#### 6.21.4.7 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.

#### 6.21.4.8 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.

#### 6.21.4.9 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.

#### 6.21.4.10 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.

#### 6.21.4.11 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.

#### 6.21.4.12 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

## 6.22 vecHasher< T > Struct Template Reference

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

#### **Public Member Functions**

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

#### 6.22.1 Detailed Description

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

Definition at line 99 of file typedefs.hpp.

#### 6.22.2 Member Function Documentation

#### 6.22.2.1 operator()()

Definition at line 100 of file typedefs.hpp.

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

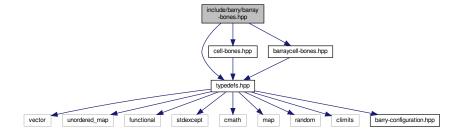
• include/barry/typedefs.hpp

## **Chapter 7**

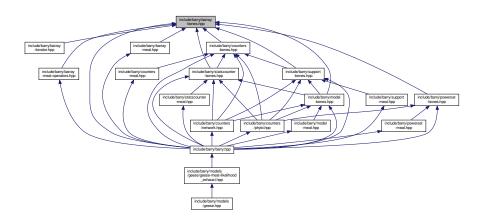
## **File Documentation**

## 7.1 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:



118 File Documentation

#### Classes

class BArray < Cell\_Type, Data\_Type >
 Baseline class for binary arrays.

#### **Macros**

• #define BARRAY\_BONES\_HPP 1

#### 7.1.1 Macro Definition Documentation

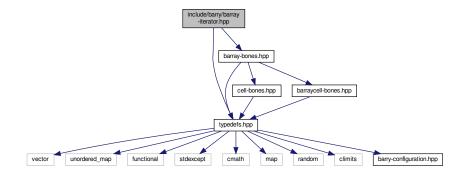
#### 7.1.1.1 BARRAY\_BONES\_HPP

```
#define BARRAY_BONES_HPP 1
```

Definition at line 8 of file barray-bones.hpp.

## 7.2 include/barry/barray-iterator.hpp File Reference

```
#include "typedefs.hpp"
#include "barray-bones.hpp"
Include dependency graph for barray-iterator.hpp:
```



#### **Classes**

 $\bullet \ \, {\sf class\ ConstBArrayRowlter}{<{\sf Cell\_Type},\ Data\_Type}>\\$ 

#### Macros

• #define BARRAY\_ITERATOR\_HPP 1

#### 7.2.1 Macro Definition Documentation

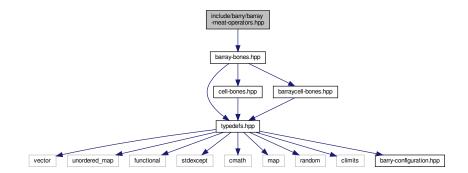
#### 7.2.1.1 BARRAY\_ITERATOR\_HPP

#define BARRAY\_ITERATOR\_HPP 1

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

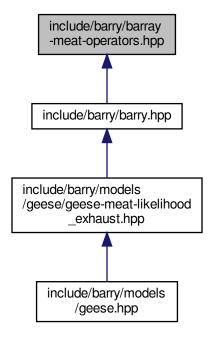
## 7.3 include/barry/barray-meat-operators.hpp File Reference

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



120 File Documentation

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)

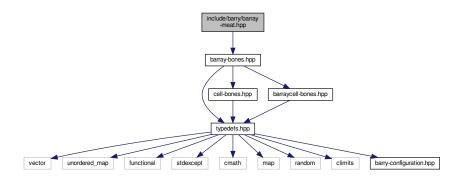
#### 7.3.1 Function Documentation

#### 7.3.1.1 checkdim\_()

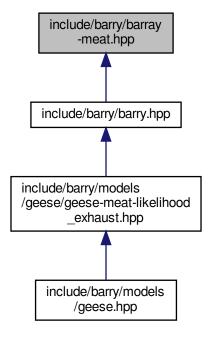
Definition at line 8 of file barray-meat-operators.hpp.

## 7.4 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:

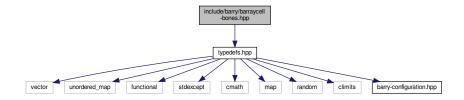


## 7.5 include/barry/barraycell-bones.hpp File Reference

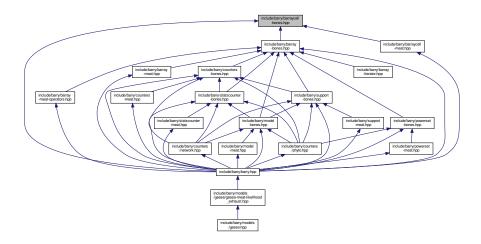
#include "typedefs.hpp"

122 File Documentation

Include dependency graph for barraycell-bones.hpp:



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

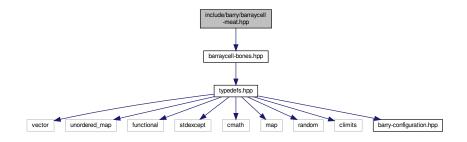


#### Classes

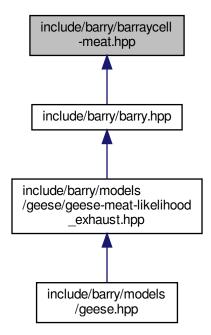
- class BArrayCell< Cell\_Type, Data\_Type>
- class BArrayCell\_const< Cell\_Type, Data\_Type >

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

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

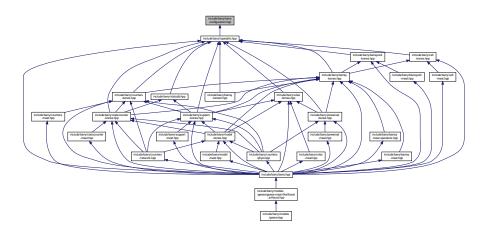


This graph shows which files directly or indirectly include this file:



## 7.7 include/barry/barry-configuration.hpp File Reference

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



### **Configuration MACROS**

These are mostly related to performance. The definitions follow:

124 File Documentation

• 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)
- template<typename Ta , typename Tb >
   using Map = std::map< Ta, Tb >

#### 7.7.1 Macro Definition Documentation

#### 7.7.1.1 BARRY\_ISFINITE

Definition at line 35 of file barry-configuration.hpp.

#### 7.7.1.2 BARRY\_SAFE\_EXP

```
#define BARRY_SAFE_EXP -100.0
```

Definition at line 28 of file barry-configuration.hpp.

#### 7.7.2 Typedef Documentation

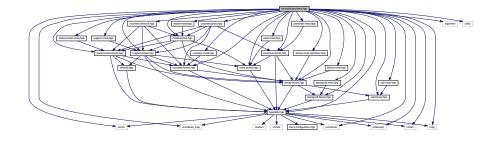
#### 7.7.2.1 Map

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

Definition at line 22 of file barry-configuration.hpp.

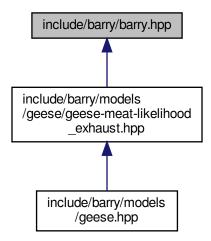
### 7.8 include/barry/barry.hpp File Reference

```
#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:
```



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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)

#### 7.8.1 Macro Definition Documentation

#### 7.8.1.1 COUNTER\_FUNCTION

#### 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.

## 7.8.1.2 COUNTER\_LAMBDA

#### 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.

## 7.8.1.3 RULE\_FUNCTION

```
#define RULE_FUNCTION(
```

#### 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.

## 7.8.1.4 RULE\_LAMBDA

```
#define RULE_LAMBDA( a )
```

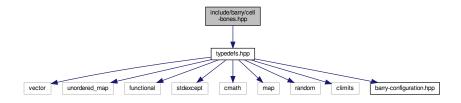
#### Value:

```
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)
```

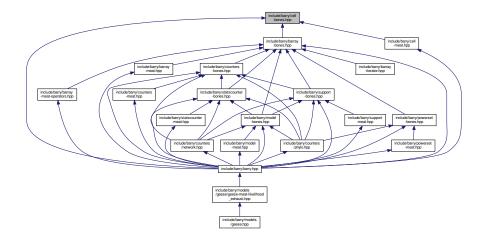
Definition at line 74 of file barry.hpp.

## 7.9 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:

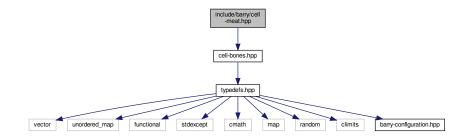


## Classes

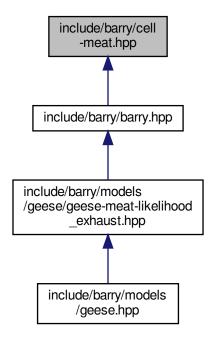
class Cell
 Cell\_Type >
 Entries in BArray. For now, it only has two members:

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

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



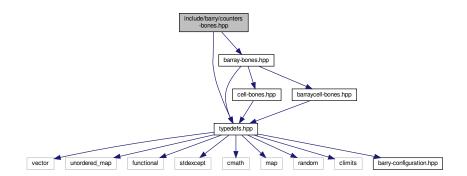
This graph shows which files directly or indirectly include this file:



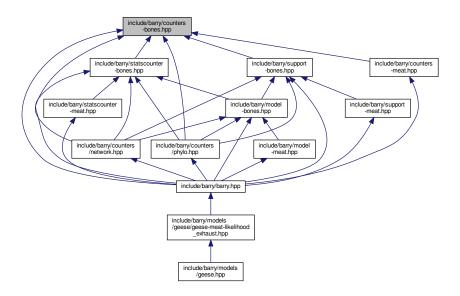
# 7.11 include/barry/col-bones.hpp File Reference

# 7.12 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:



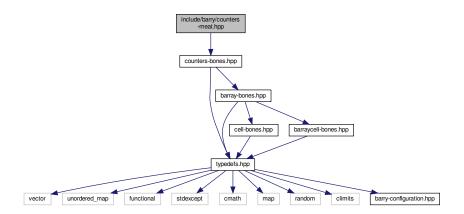
## **Classes**

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

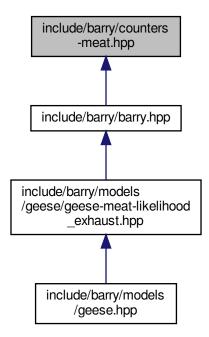
Vector of counters.

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

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



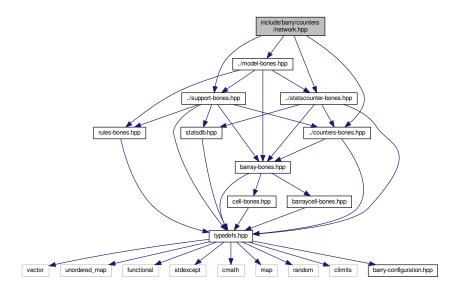
This graph shows which files directly or indirectly include this file:



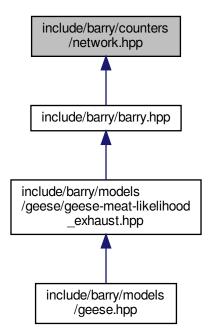
# 7.14 include/barry/counters/network.hpp File Reference

```
#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
- $\bullet \ \ typedef \ Support < Network, \ NetCounterData > NetSupport$
- typedef StatsCounter< Network, NetCounterData > NetStatsCounter
- typedef Model < Network, NetCounterData > NetModel
- typedef Rule < Network, bool > NetRule
- typedef Rules < Network, bool > NetRules

#### **Functions**

#### Counters for network models

#### Parameters

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

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.

## 7.14.1 Macro Definition Documentation

## 7.14.1.1 NET C DATA IDX

Definition at line 68 of file network.hpp.

#### 7.14.1.2 NET C DATA NUM

Definition at line 69 of file network.hpp.

#### 7.14.1.3 NETWORK\_COUNTER

Function for definition of a network counter function

Definition at line 89 of file network.hpp.

## 7.14.1.4 NETWORK\_COUNTER\_LAMBDA

Lambda function for definition of a network counter function

Definition at line 92 of file network.hpp.

## 7.14.1.5 NETWORK\_RULE

Function for definition of a network counter function

Definition at line 101 of file network.hpp.

#### 7.14.1.6 NETWORK RULE LAMBDA

Lambda function for definition of a network counter function

Definition at line 104 of file network.hpp.

## 7.14.2 Typedef Documentation

#### 7.14.2.1 NetCounter

typedef Counter<Network, NetCounterData > NetCounter

Definition at line 76 of file network.hpp.

## 7.14.2.2 NetCounters

typedef Counters< Network, NetCounterData> NetCounters

Definition at line 77 of file network.hpp.

## 7.14.2.3 NetModel

typedef Model<Network, NetCounterData> NetModel

Definition at line 80 of file network.hpp.

#### 7.14.2.4 NetRule

typedef Rule<Network,bool> NetRule

Definition at line 81 of file network.hpp.

## 7.14.2.5 NetRules

typedef Rules<Network,bool> NetRules

Definition at line 82 of file network.hpp.

## 7.14.2.6 NetStatsCounter

```
typedef StatsCounter<Network, NetCounterData> NetStatsCounter
```

Definition at line 79 of file network.hpp.

#### 7.14.2.7 NetSupport

```
typedef Support<Network, NetCounterData > NetSupport
```

Definition at line 78 of file network.hpp.

#### 7.14.2.8 Network

```
typedef BArray<double, NetworkData> Network
```

Definition at line 75 of file network.hpp.

#### 7.14.3 Function Documentation

## 7.14.3.1 counter\_absdiff()

Sum of absolute attribute difference between ego and alter.

Definition at line 405 of file network.hpp.

## 7.14.3.2 counter\_ctriads()

Definition at line 308 of file network.hpp.

## 7.14.3.3 counter\_degree()

Counts number of vertices with a given out-degree.

Definition at line 676 of file network.hpp.

## 7.14.3.4 counter\_density()

Definition at line 347 of file network.hpp.

## 7.14.3.5 counter\_diff()

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

Definition at line 447 of file network.hpp.

## 7.14.3.6 counter\_edges()

Number of edges.

Definition at line 114 of file network.hpp.

## 7.14.3.7 counter\_idegree()

Counts number of vertices with a given in-degree.

Definition at line 590 of file network.hpp.

## 7.14.3.8 counter\_idegree15()

Definition at line 363 of file network.hpp.

## 7.14.3.9 counter\_isolates()

Number of isolated vertices.

Definition at line 128 of file network.hpp.

## 7.14.3.10 counter\_istar2()

Definition at line 196 of file network.hpp.

## 7.14.3.11 counter\_mutual()

Number of mutual ties.

Definition at line 158 of file network.hpp.

## 7.14.3.12 counter\_nodecov()

Definition at line 544 of file network.hpp.

#### 7.14.3.13 counter\_nodeicov()

Definition at line 506 of file network.hpp.

## 7.14.3.14 counter\_nodematch()

Definition at line 564 of file network.hpp.

## 7.14.3.15 counter\_nodeocov()

Definition at line 525 of file network.hpp.

#### 7.14.3.16 counter odegree()

Counts number of vertices with a given out-degree.

Definition at line 632 of file network.hpp.

## 7.14.3.17 counter\_odegree15()

Definition at line 383 of file network.hpp.

## 7.14.3.18 counter\_ostar2()

Definition at line 214 of file network.hpp.

## 7.14.3.19 counter\_ttriads()

Definition at line 233 of file network.hpp.

## 7.14.3.20 NETWORK\_COUNTER()

Definition at line 489 of file network.hpp.

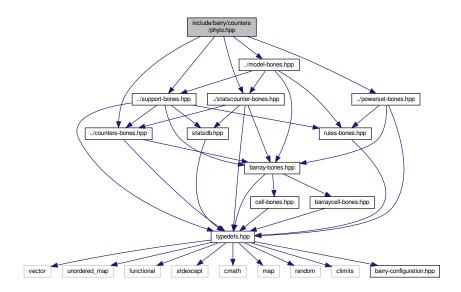
## 7.14.3.21 rules\_zerodiag()

Number of edges.

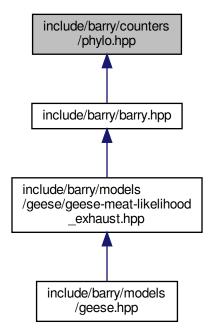
Definition at line 727 of file network.hpp.

# 7.15 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\_C\_DATA\_IDX(i) (data.operator[](i))
- #define PHYLO COUNTER(a)

Extension of a simple counter.

- #define PHYLO\_COUNTER\_LAMBDA(a)
- #define PHYLO\_CHECK\_MISSING()

## **Typedefs**

- typedef std::vector< uint > PhyloCounterData
- typedef std::vector< std::pair< uint, uint > > PhyloRuleData

#### 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**

#### Counters for phylogenetic modeling.

#### **Parameters**

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

- void counter\_overall\_gains (PhyloCounters \*counters, bool duplication=true)

  Overall functional gains.
- void counter\_gains (PhyloCounters \*counters, std::vector< uint > nfun, bool duplication=true) Functional gains for a specific function (nfun).
- void counter\_gains\_k\_offspring (PhyloCounters \*counters, std::vector< uint > nfun, uint k=1u, bool duplication=true)

Functional gains for a specific function (nfun).

- void counter\_genes\_changing (PhyloCounters \*counters, bool duplication=true)
  - Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)
- void counter\_overall\_loss (PhyloCounters \*counters, bool duplication=true)

Overall functional loss.

- void counter\_maxfuns (PhyloCounters \*counters, uint lb, uint ub, bool duplication=true)

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

  Total count of losses for an specific function.
- void counter\_overall\_changes (PhyloCounters \*counters, bool duplication=true)

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

- void counter\_subfun (PhyloCounters \*counters, uint nfunA, uint nfunB, bool duplication=true)

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

  Co-evolution (joint gain or loss)
- void counter\_longest (PhyloCounters \*counters)

Longest branch mutates (either by gain or by loss)

- void counter\_neofun (PhyloCounters \*counters, uint nfunA, uint nfunB, bool duplication=true)

  Total number of neofunctionalization events.
- void counter\_neofun\_a2b (PhyloCounters \*counters, uint nfunA, uint nfunB, bool duplication=true)

  Total number of neofunctionalization events.
- void counter\_co\_opt (PhyloCounters \*counters, uint nfunA, uint nfunB, bool duplication=true)
   Function co-opting.

## 7.15.1 Macro Definition Documentation

## 7.15.1.1 PHYLO\_C\_DATA\_IDX

Definition at line 49 of file phylo.hpp.

## 7.15.1.2 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 84 of file phylo.hpp.

## 7.15.1.3 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 78 of file phylo.hpp.

## 7.15.1.4 PHYLO\_COUNTER\_LAMBDA

Definition at line 81 of file phylo.hpp.

## 7.15.2 Typedef Documentation

## 7.15.2.1 PhyloArray

```
typedef BArray<uint, NodeData> PhyloArray
```

Definition at line 56 of file phylo.hpp.

## 7.15.2.2 PhyloCounter

typedef Counter<PhyloArray, PhyloCounterData > PhyloCounter

Definition at line 57 of file phylo.hpp.

## 7.15.2.3 PhyloCounterData

typedef std::vector< uint > PhyloCounterData

Definition at line 46 of file phylo.hpp.

## 7.15.2.4 PhyloCounters

typedef Counters< PhyloArray, PhyloCounterData> PhyloCounters

Definition at line 58 of file phylo.hpp.

## 7.15.2.5 PhyloModel

typedef Model<PhyloArray, PhyloCounterData, PhyloRuleData> PhyloModel

Definition at line 63 of file phylo.hpp.

## 7.15.2.6 PhyloPowerSet

typedef PowerSet<PhyloArray, PhyloRuleData> PhyloPowerSet

Definition at line 64 of file phylo.hpp.

## 7.15.2.7 PhyloRule

typedef Rule<PhyloArray,PhyloRuleData> PhyloRule

Definition at line 59 of file phylo.hpp.

#### 7.15.2.8 PhyloRuleData

```
typedef std::vector< std::pair< uint, uint > > PhyloRuleData
```

Definition at line 47 of file phylo.hpp.

#### 7.15.2.9 PhyloRules

```
typedef Rules<PhyloArray,PhyloRuleData> PhyloRules
```

Definition at line 60 of file phylo.hpp.

#### 7.15.2.10 PhyloStatsCounter

```
typedef StatsCounter<PhyloArray, PhyloCounterData> PhyloStatsCounter
```

Definition at line 62 of file phylo.hpp.

## 7.15.2.11 PhyloSupport

```
typedef Support<PhyloArray, PhyloCounterData, PhyloRuleData> PhyloSupport
```

Definition at line 61 of file phylo.hpp.

#### 7.15.3 Function Documentation

## 7.15.3.1 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 931 of file phylo.hpp.

#### 7.15.3.2 counter\_cogain()

Co-evolution (joint gain or loss)

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

Definition at line 618 of file phylo.hpp.

#### 7.15.3.3 counter\_gains()

Functional gains for a specific function (nfun).

Definition at line 132 of file phylo.hpp.

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

Functional gains for a specific function (nfun).

Definition at line 167 of file phylo.hpp.

#### 7.15.3.5 counter\_genes\_changing()

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

Definition at line 230 of file phylo.hpp.

## 7.15.3.6 counter\_longest()

Longest branch mutates (either by gain or by loss)

Definition at line 664 of file phylo.hpp.

#### 7.15.3.7 counter\_loss()

Total count of losses for an specific function.

Definition at line 444 of file phylo.hpp.

#### 7.15.3.8 counter\_maxfuns()

Cap the number of functions per gene.

Definition at line 371 of file phylo.hpp.

## 7.15.3.9 counter\_neofun()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 750 of file phylo.hpp.

#### 7.15.3.10 counter\_neofun\_a2b()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 823 of file phylo.hpp.

## 7.15.3.11 counter\_overall\_changes()

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

Definition at line 486 of file phylo.hpp.

## 7.15.3.12 counter\_overall\_gains()

Overall functional gains.

Total number of gains (irrespective of the function).

Definition at line 99 of file phylo.hpp.

## 7.15.3.13 counter\_overall\_loss()

Overall functional loss.

Definition at line 330 of file phylo.hpp.

## 7.15.3.14 counter\_subfun()

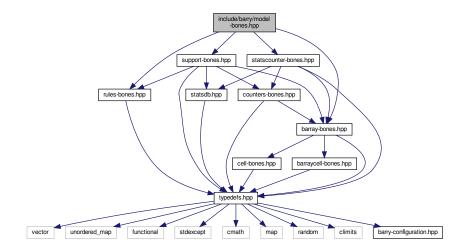
Total count of Sub-functionalization events.

It requires to specify data = {funA, funB}

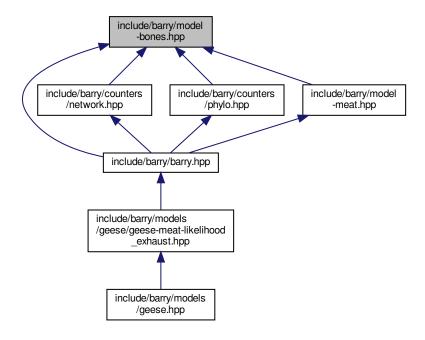
Definition at line 545 of file phylo.hpp.

## 7.16 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)

## 7.16.1 Function Documentation

#### 7.16.1.1 keygen\_default()

Array Hasher class (used for computing support)

Definition at line 63 of file model-bones.hpp.

## 7.16.1.2 likelihood\_()

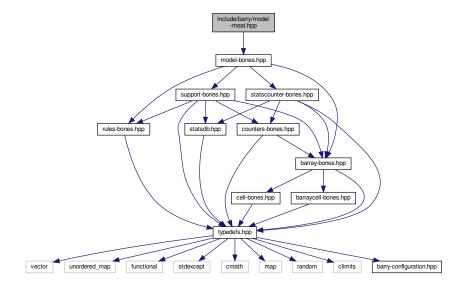
Definition at line 35 of file model-bones.hpp.

## 7.16.1.3 update\_normalizing\_constant()

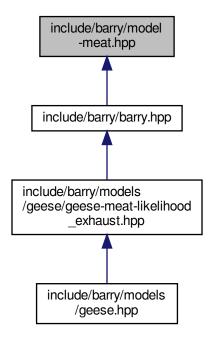
Definition at line 11 of file model-bones.hpp.

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

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

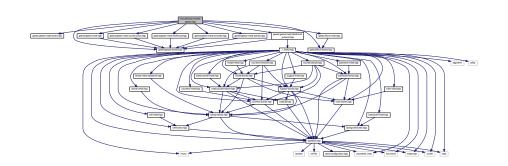


This graph shows which files directly or indirectly include this file:



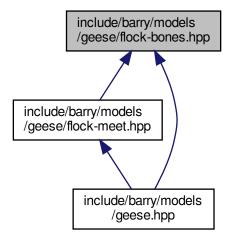
# 7.18 include/barry/models/geese.hpp File Reference

```
#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:
```



## 7.19 include/barry/models/geese/flock-bones.hpp File Reference

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



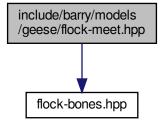
## Classes

class Flock

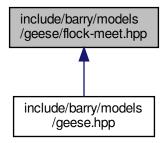
A Flock is a group of Geese.

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

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

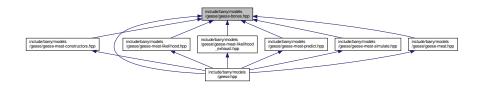


This graph shows which files directly or indirectly include this file:



## 7.21 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)

## 7.21.1 Macro Definition Documentation

#### **7.21.1.1 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.

## 7.21.2 Function Documentation

## 7.21.2.1 keygen\_full()

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

## 7.21.2.2 RULE\_FUNCTION()

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

## 7.21.2.3 vec\_diff()

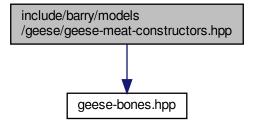
Definition at line 51 of file geese-bones.hpp.

## 7.21.2.4 vector\_caster()

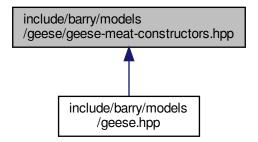
Definition at line 10 of file geese-bones.hpp.

# 7.22 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

## 7.22.1 Macro Definition Documentation

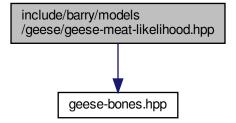
## 7.22.1.1 GEESE\_MEAT\_CONSTRUCTORS\_HPP

#define GEESE\_MEAT\_CONSTRUCTORS\_HPP 1

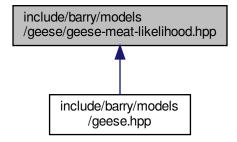
Definition at line 4 of file geese-meat-constructors.hpp.

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

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



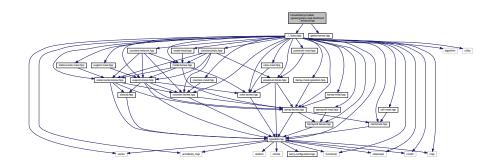
This graph shows which files directly or indirectly include this file:



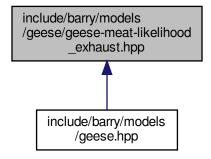
# 7.24 include/barry/models/geese/geese-meat-likelihood\_exhaust.hpp File Reference

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

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



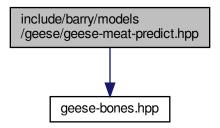
This graph shows which files directly or indirectly include this file:



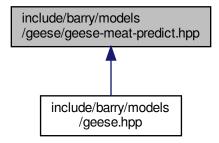
# 7.25 include/barry/models/geese/geese-meat-predict.hpp File Reference

#include "geese-bones.hpp"

Include dependency graph for geese-meat-predict.hpp:



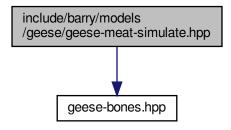
This graph shows which files directly or indirectly include this file:



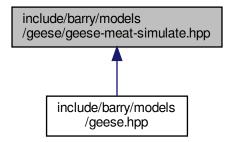
# 7.26 include/barry/models/geese/geese-meat-simulate.hpp File Reference

#include "geese-bones.hpp"

Include dependency graph for geese-meat-simulate.hpp:

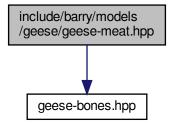


This graph shows which files directly or indirectly include this file:

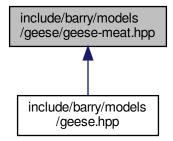


# 7.27 include/barry/models/geese/geese-meat.hpp File Reference

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

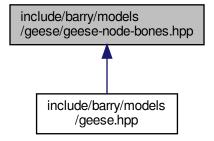


This graph shows which files directly or indirectly include this file:



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

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



#### **Classes**

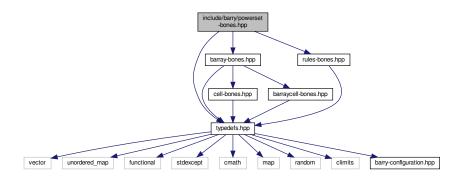
• class Node

A single node for the model.

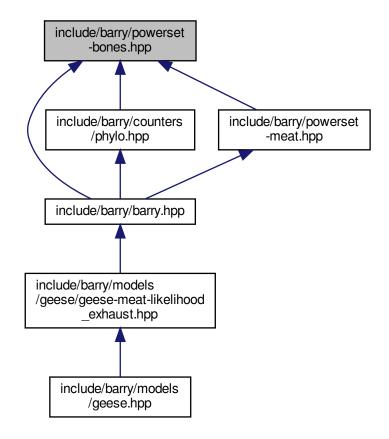
# 7.29 include/barry/powerset-bones.hpp File Reference

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

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



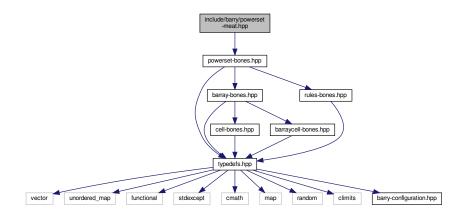
This graph shows which files directly or indirectly include this file:



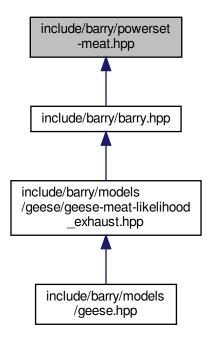
## **Classes**

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

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



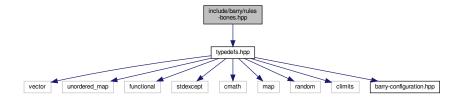
This graph shows which files directly or indirectly include this file:



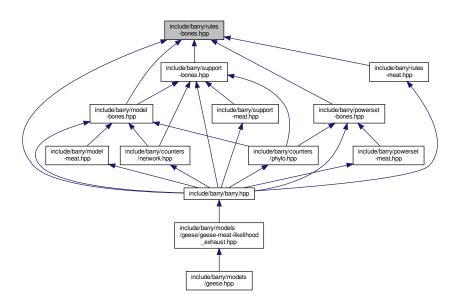
## 7.31 include/barry/rules-bones.hpp File Reference

#include "typedefs.hpp"

Include dependency graph for rules-bones.hpp:



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



#### **Classes**

- class Rule < Array\_Type, Data\_Type >
  - Rule for determining if a cell should be included in a sequence.
- class Rules < Array\_Type, Data\_Type >

Vector of objects of class Rule.

## **Functions**

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

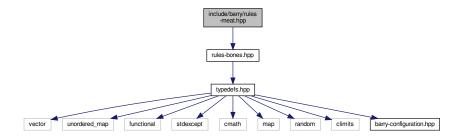
### 7.31.1 Function Documentation

#### 7.31.1.1 rule\_fun\_default()

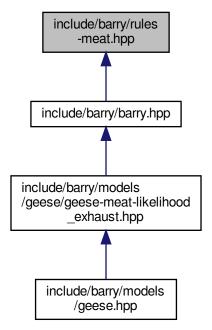
Definition at line 10 of file rules-bones.hpp.

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

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



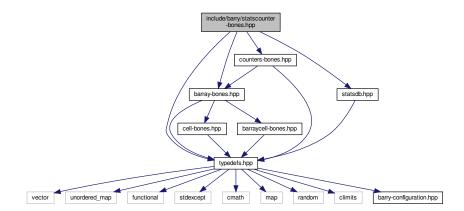
This graph shows which files directly or indirectly include this file:



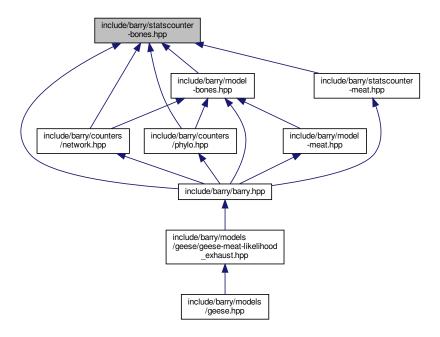
## 7.33 include/barry/statscounter-bones.hpp File Reference

```
#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:

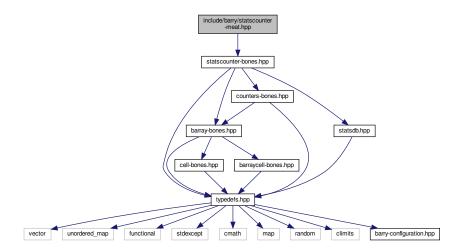


#### **Classes**

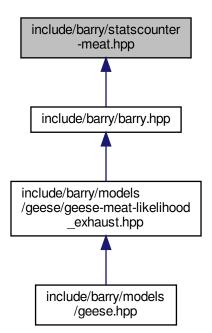
class StatsCounter < Array\_Type, Data\_Type >
 Count stats for a single Array.

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

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

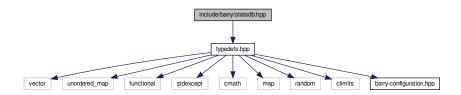


This graph shows which files directly or indirectly include this file:

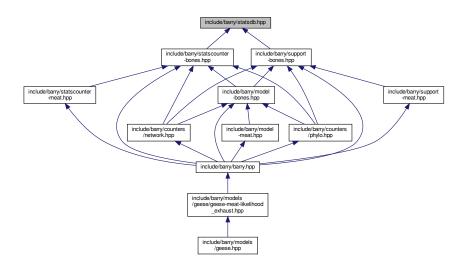


## 7.35 include/barry/statsdb.hpp File Reference

#include "typedefs.hpp"
Include dependency graph for statsdb.hpp:



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



#### **Classes**

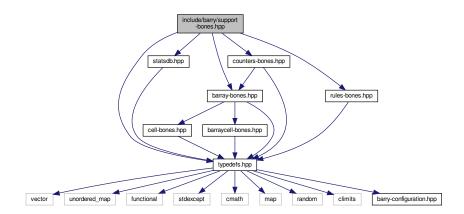
class FreqTable < T >

Database of statistics.

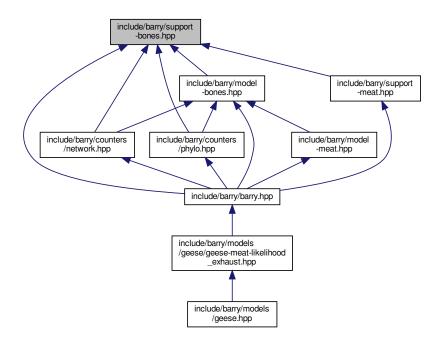
## 7.36 include/barry/support-bones.hpp File Reference

```
#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:

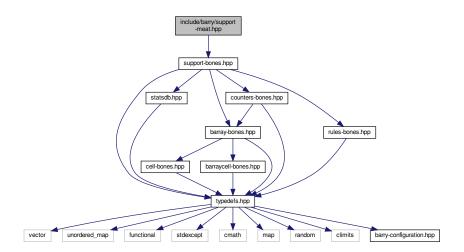


## **Classes**

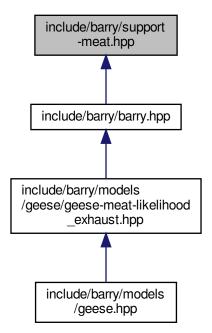
class Support < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type >
 Compute the support of sufficient statistics.

## 7.37 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

## 7.37.1 Macro Definition Documentation

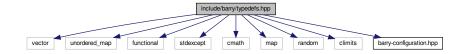
## 7.37.1.1 BARRY\_SUPPORT\_MEAT\_HPP

```
#define BARRY_SUPPORT_MEAT_HPP 1
```

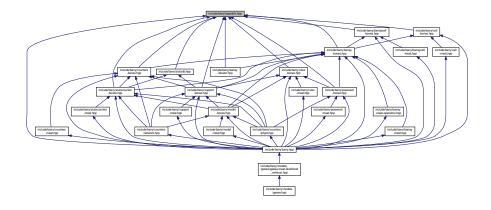
Definition at line 4 of file support-meat.hpp.

## 7.38 include/barry/typedefs.hpp File Reference

```
#include <vector>
#include <unordered_map>
#include <functional>
#include <stdexcept>
#include <cmath>
#include <map>
#include <random>
#include <climits>
#include "barry-configuration.hpp"
Include dependency graph for typedefs.hpp:
```



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



#### Classes

- class Entries < Cell\_Type >
  - A wrapper class to store source, target, val from a BArray object.
- struct vecHasher

## **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
```

#### 7.38.1 Macro Definition Documentation

## 7.38.1.1 A\_COL

Definition at line 125 of file typedefs.hpp.

#### 7.38.1.2 A\_ROW

Definition at line 124 of file typedefs.hpp.

#### 7.38.1.3 COL

Definition at line 25 of file typedefs.hpp.

#### 7.38.1.4 ROW

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

Definition at line 24 of file typedefs.hpp.

## 7.38.2 Typedef Documentation

## 7.38.2.1 Col\_type

```
template<typename Cell_Type >
using Col_type = Map< uint, Cell<Cell_Type>* >
```

Definition at line 64 of file typedefs.hpp.

#### 7.38.2.2 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 138 of file typedefs.hpp.

#### 7.38.2.3 Counts\_type

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

Definition at line 57 of file typedefs.hpp.

#### 7.38.2.4 MapVec\_type

```
template<typename Ta = double, typename Tb = uint>
using MapVec_type = std::unordered_map< std::vector< Ta >, Tb, vecHasher<Ta> >
```

Definition at line 118 of file typedefs.hpp.

#### 7.38.2.5 Row\_type

```
template<typename Cell_Type >
using Row_type = Map< uint, Cell<Cell_Type> >
```

Definition at line 61 of file typedefs.hpp.

## 7.38.2.6 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 141 of file typedefs.hpp.

#### 7.38.2.7 uint

```
typedef unsigned int uint
```

Definition at line 21 of file typedefs.hpp.

## 7.38.3 Function Documentation

### 7.38.3.1 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 152 of file typedefs.hpp.

## 7.38.3.2 vec\_equal\_approx()

Definition at line 170 of file typedefs.hpp.

#### 7.38.3.3 vec\_inner\_prod()

Definition at line 190 of file typedefs.hpp.

## 7.39 README.md File Reference

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