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. Its primary goal is to provide a general framework for building discrete exponential-family models. A particular example is Exponential Random Graph Models (ERGMs), but we can use barry to deal with non-square arrays.

Among the key features included in barry, we have:

- · Sparse arrays.
- · User-defined count statistics.
- · User-defined constrain of the support set.
- · Powerset generation of binary arrays.
- Discrete Exponential Family Models module (DEFMs).
- · Pooled DEFMs.

To use barry, you can either download the entire repository or, since it is header-only, the single header version barry.hpp.

This library was created and maintained by Dr. George G. Vega Yon as part of his doctoral dissertation "Essays on Bioinformatics and Social Network Analysis: Statistical and Computational Methods for Complex Systems."

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# **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 <ostream>
#include "../include/barry.hpp"
typedef std::vector< unsigned int > vuint;
int main() {
  // Creating network of size six with five ties
  netcounters::Network net(
       {0, 0, 4, 4, 2, 0, 1},
       {1, 2, 0, 2, 4, 0, 1}
  // How does this looks like?
  net.print("Current view");
  // Adding extra ties
  net += \{1, 0\};
  net(2, 0) = true;
  // And removing a couple
  net(0, 0) = false;
net -= {1, 1};
net.print("New view");
  // 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);
  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();
    "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
   0,]
  1,]
           1
  2,]
                   1
  3,] .
4,] 1
          . .
  5,]
New view
  0,]
           1
              1
       1
  1,]
                   1
  2,] 1
  3,] . . .
4,] 1 . 1
  5,] . .
Edges
Transitive triads : 3
Isolates
C triads
                  : 3
Mutuals
```

# **Features**

# **Efficient memory usage**

One of the key features of barry is that it will handle memory efficiently. In the case of pooled-data models, the module for statistical models avoids double-counting support when possible by keeping track of what datasets (networks, for instance) share the same.

# **Documentation**

More information can be found in the Doxygen website <a href="here">here</a> and in the PDF version of the documentation <a href="here">here</a>.

# **Code of Conduct**

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

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# **Module Index**

# 2.1 Modules

Here is a list of all modules:

Counting												 										13
Statistical Models												 										24
Network counters												 										24
Phylo counters												 										47
Phylo rules											 					 						35

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# **Hierarchical Index**

# 3.1 Class Hierarchy

This inheritance list is sorted roughly, but not completely, alphabetically:

BArray< Cell_Type, Data_Type >
BArray< bool, bool >
BArray< Cell_Type, Data_Type >
BArrayCell< Cell_Type, Data_Type >
BArrayCell_const< Cell_Type, Data_Type >
BArrayDense < Cell_Type, Data_Type >
BArrayDense < bool, bool >
BArrayDenseCell< Cell_Type, Data_Type >
BArrayDenseCell_const< Cell_Type, Data_Type >
BArrayDenseCol < Cell_Type, Data_Type >
BArrayDenseCol_const< Cell_Type, Data_Type >
BArrayDenseRow < Cell_Type, Data_Type >
BArrayDenseRow_const< Cell_Type, Data_Type >
BArrayRow< Cell_Type, Data_Type >
BArrayRow_const< Cell_Type, Data_Type >
BArrayVector< Cell_Type, Data_Type >
BArrayVector_const< Cell_Type, Data_Type >
Cell< Cell_Type >
Cell< bool >
Cell_const< Cell_Type >
ConstBArrayRowlter< Cell_Type, Data_Type >
Counter< Array_Type, Data_Type >
Counters < Array_Type, Data_Type >
Counters < Array_Type, Data_Type >
$Counters < BArray < bool, bool > \dots $
Counters < BArray <>, bool >
DEFMCounterData
DEFMData
DEFMModel
DEFM
DEFMRuleData
DEFMRuleDynData
Entries < Cell_Type >
Flock
FreqTable < T >

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Geese
Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >
NetCounterData
NetworkData
Node
NodeData
PhyloCounterData
PhyloRuleDynData
PowerSet < Array_Type, Data_Rule_Type >
Progress
Rule < Array_Type, Data_Type >
Rules < Array_Type, Data_Type >
$Rules < BArray < bool, bool > \dots $
Rules < BArray <>, bool >
StatsCounter< Array_Type, Data_Type >
StatsCounter< BArray<>, bool >
Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >
$Support < BArray <>, bool, bool > \dots $
vecHasher < T > 240

# **Class Index**

# 4.1 Class List

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

BArray< Cell_Type, Data_Type >
Baseline class for binary arrays
BArrayCell< Cell_Type, Data_Type >
BArrayCell_const< Cell_Type, Data_Type >
BArrayDense < Cell_Type, Data_Type >
Baseline class for binary arrays
BArrayDenseCell< Cell_Type, Data_Type >
BArrayDenseCell_const< Cell_Type, Data_Type >
BArrayDenseCol < Cell_Type, Data_Type >
BArrayDenseCol_const< Cell_Type, Data_Type >
BArrayDenseRow < Cell_Type, Data_Type >
BArrayDenseRow_const< Cell_Type, Data_Type >
BArrayRow < Cell_Type, Data_Type >
BArrayRow_const < Cell_Type, Data_Type >
BArrayVector< Cell_Type, Data_Type >
Row or column of a BArray
BArrayVector_const< Cell_Type, Data_Type >
Cell< Cell_Type >
Entries in BArray. For now, it only has two members:
Cell_const< Cell_Type >
ConstBArrayRowlter < Cell_Type, Data_Type >
Counter< Array_Type, Data_Type >
A counter function based on change statistics
Counters< Array_Type, Data_Type >
Vector of counters
DEFM 137
DEFMCounterData
Data class used to store arbitrary size_t or double vectors
DEFMData
Data class for DEFM arrays
DEFMRuleData
DEFMRuleDynData
Entries < Cell_Type >
A wrapper class to store source, target, val from a BArray object 146
Flock
A Flock is a group of Geese

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FreqTable < T >	
Frequency table of vectors	155
Geese	
Annotated Phylo Model	158
Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >	
General framework for discrete exponential models. This class allows generating discrete expo-	
nential models in the form of a linear exponential model:	172
NetCounterData	
Data class used to store arbitrary size_t or double vectors	194
NetworkData	
Data class for Networks	196
Node	
A single node for the model	198
NodeData	
Data definition for the PhyloArray class	204
PhyloCounterData	205
PhyloRuleDynData	208
PowerSet < Array_Type, Data_Rule_Type >	
Powerset of a binary array	210
Progress	
A simple progress bar	217
Rule < Array_Type, Data_Type >	
Rule for determining if a cell should be included in a sequence	218
Rules < Array_Type, Data_Type >	
Vector of objects of class Rule	221
StatsCounter< Array_Type, Data_Type >	
Count stats for a single Array	225
Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >	
Compute the support of sufficient statistics	229
vecHasher< T >	240

# File Index

# 5.1 File List

Here is a list of all files with brief descriptions:

include/barry/barray-bones.hpp
include/barry/barray-iterator.hpp
include/barry/barray-meat-operators.hpp
include/barry/barray-meat.hpp
include/barry/barraycell-bones.hpp
include/barry/barraycell-meat.hpp
include/barry/barraydense-bones.hpp
include/barry/barraydense-meat-operators.hpp
include/barry/barraydense-meat.hpp
include/barry/barraydensecell-bones.hpp
include/barry/barraydensecell-meat.hpp
include/barry/barraydensecol-bones.hpp
include/barry/barraydenserow-bones.hpp
include/barry/barrayrow-bones.hpp
include/barry/barrayrow-meat.hpp
include/barry/barrayvector-bones.hpp
include/barry/barrayvector-meat.hpp
include/barry/barry-configuration.hpp
include/barry/barry-debug.hpp
include/barry/barry-macros.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/freqtable.hpp
include/barry/model-bones.hpp
include/barry/model-meat.hpp
include/barry/powerset-bones.hpp
include/barry/powerset-meat.hpp
include/barry/progress.hpp
include/barry/rules-bones.hpp
include/barry/rules-meat.hpp
include/barry/statscounter-bones.hpp

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include/barry/statscounter-meat.hpp
include/barry/support-bones.hpp
include/barry/support-meat.hpp
include/barry/typedefs.hpp
include/barry/counters/network-css.hpp
$include/barry/counters/network.hpp \\ \dots \\$
$include/barry/models/defm.hpp \\ \dots \\$
include/barry/models/geese.hpp
include/barry/models/defm/counters.hpp
$include/barry/models/defm/defm-bones.hpp \\ \dots \\$
include/barry/models/defm/defm-meat.hpp
include/barry/models/defm/defm-types.hpp
$include/barry/models/defm/formula.hpp \\ \dots \\$
include/barry/models/geese/counters.hpp
include/barry/models/geese/flock-bones.hpp
include/barry/models/geese/flock-meat.hpp
include/barry/models/geese/geese-bones.hpp
$include/barry/models/geese/geese-meat-constructors. hpp \\ \dots \\$
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-predict_exhaust.hpp
include/barry/models/geese/geese-meat-predict_sim.hpp
include/barry/models/geese/geese-meat-simulate.hpp
include/barry/models/geese/geese-meat.hpp
include/barry/models/geese/geese-node-bones.hpp
include/barry/models/geese/geese-types.hpp

# **Module Documentation**

# 6.1 Counting

# **Classes**

· class NetworkData

Data class for Networks.

class Counter< Array\_Type, Data\_Type >

A counter function based on change statistics.

#### **Macros**

- #define MAKE\_DEFM\_HASHER(hasher, a, cov)
- #define MAKE\_DUPL\_VARS()
- #define IS\_EITHER() (DATA\_AT == Geese::etype\_either)
- #define IS\_DUPLICATION() ((DATA\_AT == Geese::etype\_duplication) & (DPL))
- #define IS SPECIATION() ((DATA AT == Geese::etype speciation) & (!DPL))
- #define IF\_MATCHES()
- #define IF\_NOTMATCHES()
- #define PHYLO\_COUNTER\_LAMBDA(a)

Extension of a simple counter.

- #define PHYLO\_RULE\_DYN\_LAMBDA(a)
- #define PHYLO\_CHECK\_MISSING()
- std::string get\_last\_name (size\_t d)
- void counter\_overall\_gains (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

Overall functional gains.

- void counter\_gains (PhyloCounters \*counters, std::vector < size\_t > nfun, size\_t duplication=Geese::etype\_default)

  Functional gains for a specific function (nfun).
- void counter\_gains\_k\_offspring (PhyloCounters \*counters, std::vector< size\_t > nfun, size\_t k=1u, size\_t duplication=Geese::etype\_default)

k genes gain function nfun

void counter\_genes\_changing (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

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

void counter\_preserve\_pseudogene (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_
 t duplication=Geese::etype default)

Keeps track of how many pairs of genes preserve pseudostate.

void counter\_prop\_genes\_changing (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

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

• void counter\_overall\_loss (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

Overall functional loss.

- void counter\_maxfuns (PhyloCounters \*counters, size\_t lb, size\_t ub, size\_t duplication=Geese::etype\_default)

  Cap the number of functions per gene.
- void counter\_loss (PhyloCounters \*counters, std::vector< size\_t > nfun, size\_t duplication=Geese::etype\_default)

  Total count of losses for an specific function.
- void counter\_overall\_changes (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

  Total number of changes. Use this statistic to account for "preservation".
- void counter\_subfun (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

  Total count of Sub-functionalization events.
- void counter\_cogain (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

  Co-evolution (joint gain or loss)
- void counter\_longest (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)
   Longest branch mutates (either by gain or by loss)
- void counter\_neofun (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

  Total number of neofunctionalization events.
- void counter\_pairwise\_neofun\_singlefun (PhyloCounters \*counters, size\_t nfunA, size\_t duplication=Geese::etype\_default)

  Total number of neofunctionalization events sum\_u sum\_{{w < u} [x(u,a)\*(1 x(w,a)) + (1 x(u,a)) \* x(w,a)] change stat: delta{x(u,a): 0->1} = 1 2 \* x(w,a)
- void counter\_neofun\_a2b (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

  Total number of neofunctionalization events.
- void counter\_co\_opt (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default) Function co-opting.
- void counter\_k\_genes\_changing (PhyloCounters \*counters, size\_t k, size\_t duplication=Geese::etype\_default)

  Indicator function. Equals to one if k genes changed and zero otherwise.
- void counter\_less\_than\_p\_prop\_genes\_changing (PhyloCounters \*counters, double p, size\_t duplication=Geese::etype\_default Indicator function. Equals to one if k genes changed and zero otherwise.
- void counter\_gains\_from\_0 (PhyloCounters \*counters, std::vector < size\_t > nfun, size\_t duplication=Geese::etype\_default)

  Used when all the functions are in 0 (like the root node prob.)
- void counter\_overall\_gains\_from\_0 (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

  Used when all the functions are in 0 (like the root node prob.)
- void counter\_pairwise\_overall\_change (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

  Used when all the functions are in 0 (like the root node prob.)
- void counter\_pairwise\_preserving (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_
   t duplication=Geese::etype\_default)

Used when all the functions are in 0 (like the root node prob.)

void counter\_pairwise\_first\_gain (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_←
t duplication=Geese::etype default)

Used when all the functions are in 0 (like the root node prob.)

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# 6.1.1 Detailed Description

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

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

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

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

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

#### 6.1.2 Macro Definition Documentation

#### **6.1.2.1 IF MATCHES**

```
#define IF_MATCHES( )

Value:
    MAKE_DUPL_VARS() \
    if (IS_EITHER() | IS_DUPLICATION() | IS_SPECIATION())
```

Definition at line 20 of file counters.hpp.

# 6.1.2.2 IF\_NOTMATCHES

```
#define IF_NOTMATCHES( )

Value:
    MAKE_DUPL_VARS() \
    if (!IS_EITHER() & !IS_DUPLICATION() & !IS_SPECIATION())
```

Definition at line 22 of file counters.hpp.

#### 6.1.2.3 IS\_DUPLICATION

```
#define IS_DUPLICATION( ) ((DATA_AT == Geese::etype_duplication) & (DPL))
```

Definition at line 17 of file counters.hpp.

# 6.1.2.4 IS\_EITHER

```
#define IS_EITHER( ) (DATA_AT == Geese::etype_either)
```

Definition at line 16 of file counters.hpp.

#### 6.1.2.5 IS\_SPECIATION

```
#define IS_SPECIATION( ) ((DATA_AT == Geese::etype_speciation) & (!DPL))
```

Definition at line 18 of file counters.hpp.

#### 6.1.2.6 MAKE DEFM HASHER

#### Value:

Value:

```
barry::Hasher_fun_type<DEFMArray,DEFMCounterData> \
hasher = [cov](const DEFMArray & array, DEFMCounterData * d) { \
    std::vector< double > res; \
    /* Adding the column feature */ \
    for (size_t i = 0u; i < array.nrow(); ++i) \
        res.push_back(array.D()(i, cov)); \
    /* Adding the fixed dims */ \
    for (size_t i = 0u; i < (array.nrow() - 1); ++i) \
        for (size_t j = 0u; j < array.ncol(); ++j) \
        res.push_back(array(i, j)); \
    return res; \
}</pre>
```

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

Definition at line 21 of file counters.hpp.

# 6.1.2.7 MAKE\_DUPL\_VARS

```
#define MAKE_DUPL_VARS( )
```

```
bool DPL = Array.D_ptr()->duplication; \
size_t DATA_AT = data[0u];
```

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

Definition at line 12 of file counters.hpp.

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# 6.1.2.8 PHYLO\_CHECK\_MISSING

```
#define PHYLO_CHECK_MISSING( )

Value:
    if (Array.D_ptr() == nullptr) \
    throw std::logic_error("The array data is nullptr."); \
```

Definition at line 42 of file counters.hpp.

# 6.1.2.9 PHYLO\_COUNTER\_LAMBDA

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 36 of file counters.hpp.

#### 6.1.2.10 PHYLO RULE DYN LAMBDA

Definition at line 39 of file counters.hpp.

# 6.1.3 Function Documentation

#### 6.1.3.1 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 1296 of file counters.hpp.

#### 6.1.3.2 counter\_cogain()

Co-evolution (joint gain or loss)

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

Definition at line 791 of file counters.hpp.

### 6.1.3.3 counter gains()

Functional gains for a specific function (nfun).

Definition at line 96 of file counters.hpp.

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# 6.1.3.4 counter\_gains\_from\_0()

Used when all the functions are in 0 (like the root node prob.)

Needs to specify function a.

Definition at line 1630 of file counters.hpp.

# 6.1.3.5 counter\_gains\_k\_offspring()

k genes gain function nfun

Definition at line 156 of file counters.hpp.

# 6.1.3.6 counter\_genes\_changing()

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

Definition at line 228 of file counters.hpp.

# 6.1.3.7 counter\_k\_genes\_changing()

Indicator function. Equals to one if k genes changed and zero otherwise.

Definition at line 1394 of file counters.hpp.

#### 6.1.3.8 counter\_less\_than\_p\_prop\_genes\_changing()

Indicator function. Equals to one if k genes changed and zero otherwise.

< How many genes diverge the parent

Definition at line 1514 of file counters.hpp.

# 6.1.3.9 counter\_longest()

Longest branch mutates (either by gain or by loss)

Definition at line 848 of file counters.hpp.

# 6.1.3.10 counter\_loss()

Total count of losses for an specific function.

Definition at line 591 of file counters.hpp.

# 6.1.3.11 counter\_maxfuns()

Cap the number of functions per gene.

Definition at line 529 of file counters.hpp.

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### 6.1.3.12 counter\_neofun()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1018 of file counters.hpp.

# 6.1.3.13 counter\_neofun\_a2b()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1163 of file counters.hpp.

# 6.1.3.14 counter\_overall\_changes()

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

Definition at line 643 of file counters.hpp.

# 6.1.3.15 counter\_overall\_gains()

Overall functional gains.

Total number of gains (irrespective of the function).

Definition at line 58 of file counters.hpp.

### 6.1.3.16 counter\_overall\_gains\_from\_0()

Used when all the functions are in 0 (like the root node prob.)

Needs to specify function a.

Definition at line 1696 of file counters.hpp.

# 6.1.3.17 counter\_overall\_loss()

Overall functional loss.

Definition at line 481 of file counters.hpp.

#### 6.1.3.18 counter pairwise first gain()

Used when all the functions are in 0 (like the root node prob.)

Needs to specify function a. sum  $x(a)^3(1-x(b))^3 + x(b)^3(1-x(a))^3 + x(a)^3 + x(b)^3 + (1-x(a))^3 + (1-x($ 

Definition at line 1948 of file counters.hpp.

#### 6.1.3.19 counter pairwise neofun singlefun()

Total number of neofunctionalization events sum\_u sum\_ $\{w < u\} [x(u,a)*(1 - x(w,a)) + (1 - x(u,a)) * x(w,a)] change stat: delta<math>\{x(u,a): 0->1\} = 1 - 2 * x(w,a)$ 

Definition at line 1099 of file counters.hpp.

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#### 6.1.3.20 counter\_pairwise\_overall\_change()

Used when all the functions are in 0 (like the root node prob.)

Needs to specify function a.

Definition at line 1744 of file counters.hpp.

#### 6.1.3.21 counter pairwise preserving()

Used when all the functions are in 0 (like the root node prob.)

Needs to specify function a. sum  $x(a)^3(1-x(b))^3 + x(b)^3(1-x(a))^3 + x(a)^3 + x(b)^3 + (1-x(a))^3 + (1-x($ 

Definition at line 1809 of file counters.hpp.

# 6.1.3.22 counter\_preserve\_pseudogene()

Keeps track of how many pairs of genes preserve pseudostate.

Definition at line 297 of file counters.hpp.

#### 6.1.3.23 counter\_prop\_genes\_changing()

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

Definition at line 379 of file counters.hpp.

# 6.1.3.24 counter\_subfun()

Total count of Sub-functionalization events.

It requires to specify data = {funA, funB}

Definition at line 702 of file counters.hpp.

#### 6.1.3.25 get\_last\_name()

Definition at line 45 of file counters.hpp.

# 6.2 Statistical Models

Statistical models available in barry.

### Classes

• class Model < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type >

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

class Flock

A Flock is a group of Geese.

· class Geese

Annotated Phylo Model.

# 6.2.1 Detailed Description

Statistical models available in barry.

# 6.3 Network counters

Counters for network models.

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#### **Functions**

```
• template<typename Tnet = Network>
  void counter_edges (NetCounters < Tnet > *counters)
     Number of edges.
• template<typename Tnet = Network>
  void counter isolates (NetCounters < Tnet > *counters)
     Number of isolated vertices.

    template<> void counter isolates (NetCounters< NetworkDense > *counters)

    template<typename Tnet = Network>

  void counter_mutual (NetCounters< Tnet > *counters)
     Number of mutual ties.
• template<typename Tnet = Network>
  void counter_istar2 (NetCounters < Tnet > *counters)

    template<> void counter istar2 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter ostar2 (NetCounters< Tnet > *counters)

    template<> void counter_ostar2 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter_ttriads (NetCounters < Tnet > *counters)

    template<> void counter_ttriads (NetCounters< NetworkDense > *counters)

template<typename Tnet = Network>
  void counter ctriads (NetCounters< Tnet > *counters)

    template<> void counter_ctriads (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter_density (NetCounters< Tnet > *counters)
• template<typename Tnet = Network>
  void counter_idegree15 (NetCounters < Tnet > *counters)

    template<> void counter_idegree15 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter_odegree15 (NetCounters< Tnet > *counters)

    template<> void counter_odegree15 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter_absdiff (NetCounters < Tnet > *counters, size_t attr_id, double alpha=1.0)
     Sum of absolute attribute difference between ego and alter.
• template<typename Tnet = Network>
  void counter_diff (NetCounters< Tnet > *counters, size_t 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)

• template<typename Tnet = Network>
  void counter_nodeicov (NetCounters< Tnet > *counters, size_t attr_id)
• template<typename Tnet = Network>
  void counter_nodeocov (NetCounters< Tnet > *counters, size_t attr_id)
• template<typename Tnet = Network>
  void counter_nodecov (NetCounters< Tnet > *counters, size_t attr_id)
• template<typename Tnet = Network>
  void counter_nodematch (NetCounters< Tnet > *counters, size_t attr_id)
• template<typename Tnet = Network>
  void counter idegree (NetCounters< Tnet > *counters, std::vector< size t > d)
     Counts number of vertices with a given in-degree.

    template<> void counter_idegree (NetCounters< NetworkDense > *counters, std::vector< size_t > d)

template<typename Tnet = Network>
  void counter odegree (NetCounters < Tnet > *counters, std::vector < size t > d)
     Counts number of vertices with a given out-degree.

    template<> void counter_odegree (NetCounters< NetworkDense > *counters, std::vector< size_t > d)
```

template < typename Tnet = Network>
 void counter\_degree (NetCounters < Tnet > \*counters, std::vector < size\_t > d)

Counts number of vertices with a given out-degree.

void counter\_ones (DEFMCounters \*counters, int covar\_index=-1, std::string vname="", const std::vector< std::string > \*x\_names=nullptr)

Prevalence of ones.

- void counter\_logit\_intercept (DEFMCounters \*counters, size\_t n\_y, std::vector< size\_t > which={}, int covar\_index=-1, std::string vname="", const std::vector< std::string > \*x\_names=nullptr, const std::vector< std::string > \*y\_names=nullptr)
- void counter\_transition (DEFMCounters \*counters, std::vector< size\_t > coords, std::vector< bool > signs, size\_t m\_order, size\_t n\_y, int covar\_index=-1, std::string vname="", const std::vector< std::string > \*x\_
   names=nullptr, const std::vector< std::string > \*y\_names=nullptr)

Prevalence of ones.

void counter\_transition\_formula (DEFMCounters \*counters, std::string formula, size\_t m\_order, size\_t n\_y, int covar\_index=-1, std::string vname="", const std::vector< std::string > \*x\_names=nullptr, const std::vector< std::string > \*y\_names=nullptr)

Prevalence of ones.

• void counter\_fixed\_effect (DEFMCounters \*counters, int covar\_index, double k, std::string vname="", const std::vector< std::string > \*x\_names=nullptr)

Prevalence of ones.

# Returns true if the cell is free

#### **Parameters**

rules A pointer to a DEFMRules object (Rules < DEFMArray, bool > ).

void rules\_markov\_fixed (DEFMRules \*rules, size\_t markov\_order)

Number of edges.

• void rules\_dont\_become\_zero (DEFMSupport \*support, std::vector< size\_t > ids)

Blocks switching a one to zero.

# 6.3.1 Detailed Description

Counters for network models.

#### **Parameters**

counters	A pointer to a NetCounters object (Counters <network, netcounterdata="">).</network,>
counters	A pointer to a DEFMCounters object (Counters < DEFMArray, DEFMCounterData > ).

# 6.3.2 Function Documentation

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### 6.3.2.1 counter\_absdiff()

Sum of absolute attribute difference between ego and alter.

Definition at line 908 of file network.hpp.

#### 6.3.2.2 counter\_ctriads() [1/2]

Definition at line 665 of file network.hpp.

#### 6.3.2.3 counter\_ctriads() [2/2]

Definition at line 610 of file network.hpp.

# 6.3.2.4 counter\_degree()

Counts number of vertices with a given out-degree.

Definition at line 1326 of file network.hpp.

# 6.3.2.5 counter\_density()

Definition at line 729 of file network.hpp.

# 6.3.2.6 counter\_diff()

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

Definition at line 953 of file network.hpp.

# 6.3.2.7 counter\_edges()

Number of edges.

Definition at line 152 of file network.hpp.

# 6.3.2.8 counter\_fixed\_effect()

Prevalence of ones.

#### **Parameters**

counters	Pointer ot a vector of counters
covar_index	If $>=$ than 0, then the interaction

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Definition at line 610 of file counters.hpp.

#### 6.3.2.9 counter\_idegree() [1/2]

Definition at line 1170 of file network.hpp.

#### 6.3.2.10 counter\_idegree() [2/2]

Counts number of vertices with a given in-degree.

Definition at line 1123 of file network.hpp.

# 6.3.2.11 counter\_idegree15() [1/2]

Definition at line 785 of file network.hpp.

#### 6.3.2.12 counter\_idegree15() [2/2]

Definition at line 757 of file network.hpp.

#### 6.3.2.13 counter\_isolates() [1/2]

Definition at line 215 of file network.hpp.

#### 6.3.2.14 counter\_isolates() [2/2]

Number of isolated vertices.

Definition at line 175 of file network.hpp.

#### 6.3.2.15 counter\_istar2() [1/2]

Definition at line 338 of file network.hpp.

# 6.3.2.16 counter\_istar2() [2/2]

Definition at line 312 of file network.hpp.

#### 6.3.2.17 counter\_logit\_intercept()

```
void counter_logit_intercept (
    DEFMCounters * counters,
    size_t n_y,
    std::vector< size_t > which = {},
    int covar_index = -1,
    std::string vname = "",
    const std::vector< std::string > * x_names = nullptr,
    const std::vector< std::string > * y_names = nullptr ) [inline]
```

Definition at line 151 of file counters.hpp.

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#### 6.3.2.18 counter\_mutual()

Number of mutual ties.

Definition at line 256 of file network.hpp.

#### 6.3.2.19 counter\_nodecov()

Definition at line 1066 of file network.hpp.

#### 6.3.2.20 counter\_nodeicov()

Definition at line 1016 of file network.hpp.

#### 6.3.2.21 counter\_nodematch()

Definition at line 1091 of file network.hpp.

# 6.3.2.22 counter\_nodeocov()

Definition at line 1041 of file network.hpp.

#### 6.3.2.23 counter\_odegree() [1/2]

Definition at line 1271 of file network.hpp.

# 6.3.2.24 counter\_odegree() [2/2]

Counts number of vertices with a given out-degree.

Definition at line 1223 of file network.hpp.

#### 6.3.2.25 counter\_odegree15() [1/2]

Definition at line 862 of file network.hpp.

# 6.3.2.26 counter\_odegree15() [2/2]

Definition at line 834 of file network.hpp.

#### 6.3.2.27 counter\_ones()

```
void counter_ones (
    DEFMCounters * counters,
    int covar_index = -1,
    std::string vname = "",
    const std::vector< std::string > * x_names = nullptr ) [inline]
```

Prevalence of ones.

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# **Parameters**

counters	Pointer ot a vector of counters
covar_index	If $>=$ than 0, then the interaction

Definition at line 81 of file counters.hpp.

#### 6.3.2.28 counter\_ostar2() [1/2]

Definition at line 404 of file network.hpp.

#### 6.3.2.29 counter\_ostar2() [2/2]

Definition at line 376 of file network.hpp.

# 6.3.2.30 counter\_transition()

```
void counter_transition (
    DEFMCounters * counters,
    std::vector< size_t > coords,
    std::vector< bool > signs,
    size_t m_order,
    size_t n_y,
    int covar_index = -1,
    std::string vname = "",
    const std::vector< std::string > * x_names = nullptr,
    const std::vector< std::string > * y_names = nullptr ) [inline]
```

Prevalence of ones.

#### **Parameters**

counters	Pointer ot a vector of counters
covar_index	If $>=$ than 0, then the interaction

Definition at line 270 of file counters.hpp.

#### 6.3.2.31 counter\_transition\_formula()

```
void counter_transition_formula (
    DEFMCounters * counters,
    std::string formula,
    size_t m_order,
    size_t n_y,
    int covar_index = -1,
    std::string vname = "",
    const std::vector< std::string > * x_names = nullptr,
    const std::vector< std::string > * y_names = nullptr ) [inline]
```

Prevalence of ones.

#### **Parameters**

counters	Pointer ot a vector of counters
covar_index	If $>=$ than 0, then the interaction

Definition at line 579 of file counters.hpp.

# 6.3.2.32 counter\_ttriads() [1/2]

Definition at line 531 of file network.hpp.

# 6.3.2.33 counter\_ttriads() [2/2]

Definition at line 441 of file network.hpp.

#### 6.3.2.34 NETWORK\_COUNTER()

Definition at line 997 of file network.hpp.

6.4 Phylo rules 35

# 6.3.2.35 rules\_dont\_become\_zero()

Blocks switching a one to zero.

#### **Parameters**

rules	
ids	lds of the variables that will follow this rule.

Definition at line 678 of file counters.hpp.

#### 6.3.2.36 rules\_markov\_fixed()

Number of edges.

Definition at line 653 of file counters.hpp.

# 6.4 Phylo rules

Rules for phylogenetic modeling.

Collaboration diagram for Phylo rules:



### Classes

• class DEFMRuleDynData

#### **Functions**

DEFMData::DEFMData ()

Vector indicating which covariates are included in the model.

 DEFMData::DEFMData (DEFMArray \*array\_, const double \*covariates\_, size\_t obs\_start\_, size\_t X\_ncol\_, size\_t X\_nrow\_)

Constructor.

• double DEFMData::operator() (size\_t i, size\_t j) const

Access to the row (i) colum (j) data.

- double DEFMData::at (size t i, size t j) const
- size t DEFMData::ncol () const
- size\_t DEFMData::nrow () const
- · void DEFMData::print () const
- DEFMData::~DEFMData ()
- DEFMCounterData::DEFMCounterData ()
- DEFMCounterData::DEFMCounterData (const std::vector< size\_t > indices\_, const std::vector< double > numbers\_, const std::vector< bool > logical\_, bool is\_motif\_=true)
- size\_t DEFMCounterData::idx (size\_t i) const
- double DEFMCounterData::num (size\_t i) const
- bool DEFMCounterData::is\_true (size\_t i) const
- DEFMCounterData::~DEFMCounterData ()
- double DEFMRuleData::num (size\_t i) const
- size\_t DEFMRuleData::idx (size\_t i) const
- bool DEFMRuleData::is\_true (size\_t i) const
- DEFMRuleData::DEFMRuleData ()
- DEFMRuleData::DEFMRuleData (std::vector< double > numbers\_, std::vector< size\_t > indices\_, std
   ::vector< bool > logical\_)
- DEFMRuleData::DEFMRuleData (std::vector< double > numbers\_, std::vector< size\_t > indices\_)
- DEFMRuleDynData::DEFMRuleDynData (const std::vector< double > \*counts\_, std::vector< double > numbers\_={}, std::vector< size\_t > indices\_={}, std::vector< bool > logical\_={})
- DEFMRuleDynData::~DEFMRuleDynData ()
- void rule\_dyn\_limit\_changes (PhyloSupport \*support, size\_t pos, size\_t lb, size\_t ub, size\_← t duplication=Geese::etype\_default)

Overall functional gains.

### **Variables**

- DEFMArray \* DEFMData::array
- const double \* DEFMData::covariates

Vector of covariates (complete vector)

· size\_t DEFMData::obs\_start

Index of the observation in the data.

size\_t DEFMData::X\_ncol

Number of columns in the array of covariates.

size\_t DEFMData::X\_nrow

Number of rows in the array of covariates.

- std::vector< size\_t > DEFMData::covar\_sort
- std::vector< size\_t > DEFMData::covar\_used

Value where the sorting of the covariates is stored.

- std::vector< size\_t > DEFMCounterData::indices
- std::vector< double > DEFMCounterData::numbers
- std::vector< bool > DEFMCounterData::logical
- · bool DEFMCounterData::is motif

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If false, then is a logit intercept.

- std::vector< double > DEFMRuleData::numbers
- std::vector< size t > DEFMRuleData::indices
- std::vector< bool > DEFMRuleData::logical
- bool DEFMRuleData::init = false
- const std::vector< double > \* DEFMRuleDynData::counts

## Convenient typedefs for network objects.

- typedef barry::Counter
   DEFMArray, DEFMCounterData > DEFMCounter
- typedef barry::Counters < DEFMCounterData > DEFMCounters
- typedef barry::Support
   DEFMArray, DEFMCounterData, DEFMRuleData, DEFMRuleDynData > DEFMSupport
- typedef barry::StatsCounter
   DEFMArray, DEFMCounterData > DEFMStatsCounter
- typedef barry::Model< DEFMArray, DEFMCounterData, DEFMRuleData, DEFMRuleDynData > DEFMModel
- typedef barry::Rule < DEFMArray, DEFMRuleData > DEFMRule
- typedef barry::Rules< DEFMArray, DEFMRuleData > DEFMRules
- typedef barry::Rule < DEFMArray, DEFMRuleDynData > DEFMRuleDyn
- typedef barry::Rules < DEFMArray, DEFMRuleDynData > DEFMRulesDyn

#### 6.4.1 Detailed Description

Rules for phylogenetic modeling.

#### **Parameters**

rules A pointer to a PhyloRules object (Rules < PhyloArray, PhyloRuleData > ).

#### 6.4.2 Typedef Documentation

#### 6.4.2.1 DEFMCounter

typedef barry::Counter<DEFMArray, DEFMCounterData > DEFMCounter

Definition at line 171 of file defm-types.hpp.

## 6.4.2.2 DEFMCounters

typedef barry::Counters<DEFMArray, DEFMCounterData> DEFMCounters

Definition at line 172 of file defm-types.hpp.

#### 6.4.2.3 DEFMModel

typedef barry::Model<DEFMArray, DEFMCounterData,DEFMRuleData,DEFMRuleDynData> DEFMModel

Definition at line 175 of file defm-types.hpp.

#### 6.4.2.4 DEFMRule

```
typedef barry::Rule<DEFMArray, DEFMRuleData> DEFMRule
```

Definition at line 178 of file defm-types.hpp.

#### 6.4.2.5 DEFMRuleDyn

```
typedef barry::Rule<DEFMArray, DEFMRuleDynData> DEFMRuleDyn
```

Definition at line 180 of file defm-types.hpp.

#### 6.4.2.6 DEFMRules

```
typedef barry::Rules<DEFMArray, DEFMRuleData> DEFMRules
```

Definition at line 179 of file defm-types.hpp.

#### 6.4.2.7 DEFMRulesDyn

```
typedef barry::Rules<DEFMArray, DEFMRuleDynData> DEFMRulesDyn
```

Definition at line 181 of file defm-types.hpp.

#### 6.4.2.8 DEFMStatsCounter

typedef barry::StatsCounter<DEFMArray, DEFMCounterData> DEFMStatsCounter

Definition at line 174 of file defm-types.hpp.

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#### 6.4.2.9 DEFMSupport

typedef barry::Support<DEFMArray, DEFMCounterData, DEFMRuleData,DEFMRuleDynData> DEFMSupport
Definition at line 173 of file defm-types.hpp.

#### 6.4.3 Function Documentation

#### 6.4.3.1 at()

```
double DEFMData::at (  \mbox{size\_t} \ i, \\ \mbox{size\_t} \ j \mbox{) const}
```

#### 6.4.3.2 DEFMCounterData() [1/2]

```
DEFMCounterData::DEFMCounterData ( ) [inline]
```

Definition at line 72 of file defm-types.hpp.

#### 6.4.3.3 DEFMCounterData() [2/2]

```
DEFMCounterData::DEFMCounterData (
    const std::vector< size_t > indices_,
    const std::vector< double > numbers_,
    const std::vector< bool > logical_,
    bool is_motif_ = true ) [inline]
```

Definition at line 73 of file defm-types.hpp.

#### 6.4.3.4 **DEFMData()** [1/2]

```
DEFMData::DEFMData ( ) [inline]
```

Vector indicating which covariates are included in the model.

Definition at line 27 of file defm-types.hpp.

#### 6.4.3.5 **DEFMData()** [2/2]

Constructor.

#### **Parameters**

Pointer to the attribute data.
Location of the current observation in the covariates vector
Number of columns (covariates.)

Definition at line 36 of file defm-types.hpp.

# 6.4.3.6 **DEFMRuleData()** [1/3]

```
DEFMRuleData::DEFMRuleData ( ) [inline]
```

Definition at line 102 of file defm-types.hpp.

#### 6.4.3.7 **DEFMRuleData()** [2/3]

Definition at line 110 of file defm-types.hpp.

#### 6.4.3.8 **DEFMRuleData()** [3/3]

```
DEFMRuleData::DEFMRuleData (
          std::vector< double > numbers_,
          std::vector< size_t > indices_,
          std::vector< bool > logical_ ) [inline]
```

Definition at line 104 of file defm-types.hpp.

#### 6.4.3.9 DEFMRuleDynData()

```
DEFMRuleDynData::DEFMRuleDynData (
    const std::vector< double > * counts_,
    std::vector< double > numbers_ = {},
    std::vector< size_t > indices_ = {},
    std::vector< bool > logical_ = {} ) [inline]
```

Definition at line 156 of file defm-types.hpp.

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#### 6.4.3.10 idx() [1/2]

Definition at line 81 of file defm-types.hpp.

# 6.4.3.11 idx() [2/2]

Definition at line 99 of file defm-types.hpp.

### 6.4.3.12 is\_true() [1/2]

Definition at line 83 of file defm-types.hpp.

#### 6.4.3.13 is\_true() [2/2]

Definition at line 100 of file defm-types.hpp.

#### 6.4.3.14 ncol()

```
size_t DEFMData::ncol ( ) const [inline]
```

Definition at line 123 of file defm-types.hpp.

## 6.4.3.15 nrow()

```
size_t DEFMData::nrow ( ) const [inline]
```

Definition at line 127 of file defm-types.hpp.

# 6.4.3.16 num() [1/2]

```
double DEFMCounterData::num ( \label{eq:const} \texttt{size\_t} \ i \ ) \ \texttt{const} \ \ [\texttt{inline}]
```

Definition at line 82 of file defm-types.hpp.

# 6.4.3.17 num() [2/2]

Definition at line 98 of file defm-types.hpp.

### 6.4.3.18 operator()()

Access to the row (i) colum (j) data.

# **Parameters**



Returns

double

Definition at line 118 of file defm-types.hpp.

# 6.4.3.19 print()

```
void DEFMData::print ( ) const [inline]
```

Definition at line 131 of file defm-types.hpp.

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# 6.4.3.20 rule\_dyn\_limit\_changes()

Overall functional gains.

#### **Parameters**

support	Support of a model.
pos	Position of the focal statistic.
lb	Lower bound
ub	Upper bound

#### Returns

(void) adds a rule limiting the support of the model.

Definition at line 2058 of file counters.hpp.

#### 6.4.3.21 ∼DEFMCounterData()

```
{\tt DEFMCounterData::}{\sim}{\tt DEFMCounterData~(~)} \quad [{\tt inline}]
```

Definition at line 85 of file defm-types.hpp.

#### 6.4.3.22 ∼DEFMData()

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

Definition at line 58 of file defm-types.hpp.

#### 6.4.3.23 ~DEFMRuleDynData()

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

Definition at line 163 of file defm-types.hpp.

# 6.4.4 Variable Documentation

#### 6.4.4.1 array

```
DEFMArray* DEFMData::array
```

Definition at line 19 of file defm-types.hpp.

#### 6.4.4.2 counts

```
const std::vector< double >* DEFMRuleDynData::counts
```

Definition at line 154 of file defm-types.hpp.

#### 6.4.4.3 covar\_sort

```
std::vector< size_t > DEFMData::covar_sort
```

Definition at line 24 of file defm-types.hpp.

# 6.4.4.4 covar\_used

```
std::vector< size_t > DEFMData::covar_used
```

Value where the sorting of the covariates is stored.

Definition at line 25 of file defm-types.hpp.

#### 6.4.4.5 covariates

```
const double* DEFMData::covariates
```

Vector of covariates (complete vector)

Definition at line 20 of file defm-types.hpp.

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#### 6.4.4.6 indices [1/2]

```
std::vector< size_t > DEFMCounterData::indices
```

Definition at line 67 of file defm-types.hpp.

# 6.4.4.7 indices [2/2]

```
std::vector< size_t > DEFMRuleData::indices
```

Definition at line 93 of file defm-types.hpp.

#### 6.4.4.8 init

```
bool DEFMRuleData::init = false
```

Definition at line 96 of file defm-types.hpp.

# 6.4.4.9 is\_motif

bool DEFMCounterData::is\_motif

If false, then is a logit intercept.

Definition at line 70 of file defm-types.hpp.

#### 6.4.4.10 logical [1/2]

```
std::vector< bool > DEFMCounterData::logical
```

Definition at line 69 of file defm-types.hpp.

# 6.4.4.11 logical [2/2]

std::vector< bool > DEFMRuleData::logical

Definition at line 94 of file defm-types.hpp.

#### 6.4.4.12 numbers [1/2]

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

Definition at line 68 of file defm-types.hpp.

#### 6.4.4.13 numbers [2/2]

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

Definition at line 92 of file defm-types.hpp.

# 6.4.4.14 obs\_start

```
size_t DEFMData::obs_start
```

Index of the observation in the data.

Definition at line 21 of file defm-types.hpp.

# 6.4.4.15 X\_ncol

```
size_t DEFMData::X_ncol
```

Number of columns in the array of covariates.

Definition at line 22 of file defm-types.hpp.

# 6.4.4.16 X\_nrow

```
size_t DEFMData::X_nrow
```

Number of rows in the array of covariates.

Definition at line 23 of file defm-types.hpp.

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# 6.5 Phylo counters

Counters for phylogenetic modeling.

Collaboration diagram for Phylo counters:



#### **Modules**

· Phylo rules

Rules for phylogenetic modeling.

- void counter\_overall\_gains (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)
   Overall functional gains.
- void counter\_gains (PhyloCounters \*counters, std::vector < size\_t > nfun, size\_t duplication=Geese::etype\_default)

  Functional gains for a specific function (nfun).
- void counter\_gains\_k\_offspring (PhyloCounters \*counters, std::vector< size\_t > nfun, size\_t k=1u, size\_t duplication=Geese::etype\_default)

k genes gain function nfun

- void counter\_genes\_changing (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)
  - Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)
- void counter\_preserve\_pseudogene (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_
   t duplication=Geese::etype\_default)

Keeps track of how many pairs of genes preserve pseudostate.

- void counter\_prop\_genes\_changing (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)
  - Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)
- void counter\_overall\_loss (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default) Overall functional loss.
- void counter\_maxfuns (PhyloCounters \*counters, size\_t lb, size\_t duplication=Geese::etype\_default)

  Cap the number of functions per gene.
- void counter\_loss (PhyloCounters \*counters, std::vector < size\_t > nfun, size\_t duplication=Geese::etype\_default)

  Total count of losses for an specific function.
- void counter\_overall\_changes (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)
   Total number of changes. Use this statistic to account for "preservation".
- void counter\_subfun (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

  Total count of Sub-functionalization events.
- void counter\_cogain (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

  Co-evolution (joint gain or loss)
- $\bullet \ \ void\ counter\_longest\ (PhyloCounters\ *counters,\ size\_t\ duplication=Geese::etype\_default)$ 
  - Longest branch mutates (either by gain or by loss)
- void counter\_neofun (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

  Total number of neofunctionalization events.

• void counter\_pairwise\_neofun\_singlefun (PhyloCounters \*counters, size\_t nfunA, size\_t duplication=Geese::etype\_default)

Total number of neofunctionalization events sum\_u sum\_{{w < u} [x(u,a)\*(1 - x(w,a)) + (1 - x(u,a)) \* x(w,a)] change stat: delta{x(u,a): 0->1} = 1 - 2 \* x(w,a)

- void counter\_neofun\_a2b (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

  Total number of neofunctionalization events.
- void counter\_co\_opt (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

  Function co-opting.
- void counter\_k\_genes\_changing (PhyloCounters \*counters, size\_t k, size\_t duplication=Geese::etype\_default)

  Indicator function. Equals to one if k genes changed and zero otherwise.
- void counter\_less\_than\_p\_prop\_genes\_changing (PhyloCounters \*counters, double p, size\_t duplication=Geese::etype\_default Indicator function. Equals to one if k genes changed and zero otherwise.
- void counter\_gains\_from\_0 (PhyloCounters \*counters, std::vector < size\_t > nfun, size\_t duplication=Geese::etype\_default)

  Used when all the functions are in 0 (like the root node prob.)
- void counter\_overall\_gains\_from\_0 (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

  Used when all the functions are in 0 (like the root node prob.)
- void counter\_pairwise\_overall\_change (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

  Used when all the functions are in 0 (like the root node prob.)
- void counter\_pairwise\_preserving (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_ t duplication=Geese::etype\_default)

Used when all the functions are in 0 (like the root node prob.)

void counter\_pairwise\_first\_gain (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_
 t duplication=Geese::etype\_default)

Used when all the functions are in 0 (like the root node prob.)

#### 6.5.1 Detailed Description

Counters for phylogenetic modeling.

**Parameters** 

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

### 6.5.2 Function Documentation

#### 6.5.2.1 counter\_co\_opt()

Function co-opting.

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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 1296 of file counters.hpp.

#### 6.5.2.2 counter\_cogain()

Co-evolution (joint gain or loss)

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

Definition at line 791 of file counters.hpp.

### 6.5.2.3 counter\_gains()

Functional gains for a specific function (nfun).

Definition at line 96 of file counters.hpp.

### 6.5.2.4 counter\_gains\_from\_0()

Used when all the functions are in 0 (like the root node prob.)

Needs to specify function a.

Definition at line 1630 of file counters.hpp.

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

k genes gain function nfun

Definition at line 156 of file counters.hpp.

#### 6.5.2.6 counter\_genes\_changing()

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

Definition at line 228 of file counters.hpp.

#### 6.5.2.7 counter\_k\_genes\_changing()

Indicator function. Equals to one if k genes changed and zero otherwise.

Definition at line 1394 of file counters.hpp.

# 6.5.2.8 counter\_less\_than\_p\_prop\_genes\_changing()

Indicator function. Equals to one if k genes changed and zero otherwise.

< How many genes diverge the parent

Definition at line 1514 of file counters.hpp.

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#### 6.5.2.9 counter\_longest()

Longest branch mutates (either by gain or by loss)

Definition at line 848 of file counters.hpp.

#### 6.5.2.10 counter\_loss()

Total count of losses for an specific function.

Definition at line 591 of file counters.hpp.

#### 6.5.2.11 counter\_maxfuns()

Cap the number of functions per gene.

Definition at line 529 of file counters.hpp.

# 6.5.2.12 counter\_neofun()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1018 of file counters.hpp.

#### 6.5.2.13 counter\_neofun\_a2b()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1163 of file counters.hpp.

#### 6.5.2.14 counter\_overall\_changes()

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

Definition at line 643 of file counters.hpp.

# 6.5.2.15 counter\_overall\_gains()

Overall functional gains.

Total number of gains (irrespective of the function).

Definition at line 58 of file counters.hpp.

# 6.5.2.16 counter\_overall\_gains\_from\_0()

Used when all the functions are in 0 (like the root node prob.)

Needs to specify function a.

Definition at line 1696 of file counters.hpp.

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#### 6.5.2.17 counter\_overall\_loss()

Overall functional loss.

Definition at line 481 of file counters.hpp.

#### 6.5.2.18 counter\_pairwise\_first\_gain()

Used when all the functions are in 0 (like the root node prob.)

Needs to specify function a. sum  $x(a)^3(1-x(b))^3 + x(b)^3(1-x(a))^3 + x(a)^3 + x(b)^3 + (1-x(a))^3 + (1-x($ 

Definition at line 1948 of file counters.hpp.

# 6.5.2.19 counter\_pairwise\_neofun\_singlefun()

Total number of neofunctionalization events sum\_u sum\_{w < u} [x(u,a)\*(1 - x(w,a)) + (1 - x(u,a)) \* x(w,a)] change stat: delta{x(u,a): 0->1} = 1 - 2 \* x(w,a)

Definition at line 1099 of file counters.hpp.

#### 6.5.2.20 counter\_pairwise\_overall\_change()

Used when all the functions are in 0 (like the root node prob.)

Needs to specify function a.

Definition at line 1744 of file counters.hpp.

#### 6.5.2.21 counter\_pairwise\_preserving()

Used when all the functions are in 0 (like the root node prob.)

Needs to specify function a. sum  $x(a)^3(1-x(b))^3 + x(b)^3(1-x(a))^3 + x(a)^3 + x(b)^3 + (1-x(a))^3 + (1-x($ 

Definition at line 1809 of file counters.hpp.

#### 6.5.2.22 counter\_preserve\_pseudogene()

Keeps track of how many pairs of genes preserve pseudostate.

Definition at line 297 of file counters.hpp.

### 6.5.2.23 counter\_prop\_genes\_changing()

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

Definition at line 379 of file counters.hpp.

#### 6.5.2.24 counter\_subfun()

Total count of Sub-functionalization events.

It requires to specify data = {funA, funB}

Definition at line 702 of file counters.hpp.

# **Chapter 7**

# **Namespace Documentation**

# 7.1 barry Namespace Reference

barry: Your go-to motif accountant

# **Namespaces**

counters

Tree class and Treelterator class.

# 7.1.1 Detailed Description

barry: Your go-to motif accountant

# 7.2 barry::counters Namespace Reference

Tree class and Treelterator class.

# **Namespaces**

network

# 7.2.1 Detailed Description

Tree class and Treelterator class.

# 7.3 barry::counters::network Namespace Reference

# 7.4 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

# 7.4.1 Detailed Description

Integer constants used to specify which cell should be check.

#### 7.4.2 Variable Documentation

#### 7.4.2.1 BOTH

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

Definition at line 27 of file typedefs.hpp.

# 7.4.2.2 NONE

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

Definition at line 28 of file typedefs.hpp.

#### 7.4.2.3 ONE

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

Definition at line 29 of file typedefs.hpp.

# 7.4.2.4 TWO

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

Definition at line 30 of file typedefs.hpp.

# 7.5 defm Namespace Reference

# 7.6 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

# 7.6.1 Detailed Description

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

# 7.6.2 Variable Documentation

# 7.6.2.1 AS\_ONE

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

Definition at line 45 of file typedefs.hpp.

# 7.6.2.2 AS\_ZERO

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

Definition at line 44 of file typedefs.hpp.

# 7.6.2.3 BOTH

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

Definition at line 38 of file typedefs.hpp.

#### 7.6.2.4 NONE

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

Definition at line 39 of file typedefs.hpp.

#### 7.6.2.5 ONE

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

Definition at line 40 of file typedefs.hpp.

# 7.6.2.6 TWO

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

Definition at line 41 of file typedefs.hpp.

#### 7.6.2.7 UKNOWN

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

Definition at line 43 of file typedefs.hpp.

# 7.7 geese Namespace Reference

# **Chapter 8**

# **Class Documentation**

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

Baseline class for binary arrays.

#include <barray-bones.hpp>

#### **Public Member Functions**

- bool operator== (const BArray< Cell\_Type, Data\_Type > &Array\_)
- $\sim$ BArray ()
- void out\_of\_range (size\_t i, size\_t j) const
- Cell\_Type get\_cell (size\_t i, size\_t j, bool check\_bounds=true) const
- std::vector< Cell\_Type > get\_col\_vec (size\_t i, bool check\_bounds=true) const
- std::vector< Cell Type > get row vec (size t i, bool check bounds=true) const
- void get\_col\_vec (std::vector< Cell\_Type > \*x, size\_t i, bool check\_bounds=true) const
- void get\_row\_vec (std::vector< Cell\_Type > \*x, size\_t i, bool check\_bounds=true) const
- const Row\_type< Cell\_Type > & row (size\_t i, bool check\_bounds=true) const
- const Col\_type< Cell\_Type > & col (size\_t i, bool check\_bounds=true) const
- Entries < Cell\_Type > get\_entries () const

#### Get the edgelist.

- void transpose ()
- void clear (bool hard=true)
- void resize (size\_t N\_, size\_t M\_)
- void reserve ()
- void print (const char \*fmt=nullptr,...) const
- bool is\_dense () const noexcept

#### Constructors

#### **Parameters**

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

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• BArray ()

Zero-size array.

BArray (size\_t N\_, size\_t M\_)

Empty array.

BArray (size\_t N\_, size\_t M\_, const std::vector< size\_t > &source, const std::vector< size\_t > &target, const std::vector< Cell\_Type > &value, bool add=true)

Edgelist with data.

BArray (size\_t N\_, size\_t M\_, const std::vector< size\_t > &source, const std::vector< size\_t > &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.
- void set\_data (Data\_Type \*data\_, bool delete\_data\_=false)

Set the data object.

- Data Type \* D ptr ()
- const Data Type \* D ptr () const
- Data\_Type & D ()
- const Data\_Type & D () const
- void flush\_data ()

### 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 (size\_t i, size\_t j, bool check\_bounds=true) const
- size\_t nrow () const noexcept
- size\_t ncol () const noexcept
- size\_t nnozero () const noexcept
- Cell< Cell\_Type > default\_val () 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	Wither check if the cell exists (before trying to delete/add), or, in the case of swap_cells, check if either of both cells exists/don't exist.

- BArray< Cell\_Type, Data\_Type > & operator+= (const std::pair< size\_t, size\_t > &coords)
- BArray< Cell\_Type, Data\_Type > & operator-= (const std::pair< size\_t, size\_t > &coords)
- BArrayCell< Cell\_Type, Data\_Type > operator() (size\_t i, size\_t j, bool check\_bounds=true)
- const Cell\_Type operator() (size\_t i, size\_t j, bool check\_bounds=true) const
- void rm\_cell (size\_t i, size\_t j, bool check\_bounds=true, bool check\_exists=true)
- void insert\_cell (size\_t i, size\_t j, const Cell< Cell\_Type > &v, bool check\_bounds, bool check\_exists)
- void insert cell (size ti, size ti, Cell< Cell Type > &&v, bool check bounds, bool check exists)
- void insert\_cell (size\_t i, size\_t j, Cell\_Type v, bool check\_bounds, bool check\_exists)
- void swap\_cells (size\_t i0, size\_t j0, size\_t i1, size\_t j1, bool check\_bounds=true, int check\_exists=CHECK::BOTH, int \*report=nullptr)
- void toggle cell (size t i, size t j, bool check bounds=true, int check exists=EXISTS::UKNOWN)
- void toggle\_lock (size\_t i, size\_t j, bool check\_bounds=true)

#### Column/row wise interchange

- void swap rows (size t i0, size t i1, bool check bounds=true)
- void swap cols (size t j0, size t j1, bool check bounds=true)
- void zero row (size t i, bool check bounds=true)
- void zero col (size t 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**

· bool visited = false

#### **Friends**

- class BArrayCell
   Cell Type, Data Type
- class BArrayCell const< Cell Type, Data Type >

#### 8.1.1 Detailed Description

 $\label{template} $$ \ensuremath{\sf template}$$ < typename Cell_Type = bool, typename Data_Type = bool> $$ \ensuremath{\sf class BArray}$ < Cell_Type, Data_Type> $$$ 

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.  $std::vector < std::unordered_maps < size_t, Cell_Type > >$ .

#### **Template Parameters**

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

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

# 8.1.2 Constructor & Destructor Documentation

# 8.1.2.1 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 69 of file barray-bones.hpp.

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

Empty array.

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

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

Edgelist with data.

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

Edgelist with no data (simpler)

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

Copy constructor.

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

Move operator.

#### 8.1.2.7 ∼BArray()

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

# 8.1.3 Member Function Documentation

## 8.1.3.1 clear()

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#### 8.1.3.2 col()

#### 8.1.3.3 D() [1/2]

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

#### 8.1.3.4 D() [2/2]

```
template<typename Cell_Type = bool, typename Data_Type = bool>
const Data_Type& BArray< Cell_Type, Data_Type >::D ( ) const
```

#### 8.1.3.5 D\_ptr() [1/2]

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

# 8.1.3.6 D\_ptr() [2/2]

```
template<typename Cell_Type = bool, typename Data_Type = bool>
const Data_Type* BArray< Cell_Type, Data_Type >::D_ptr () const
```

# 8.1.3.7 default\_val()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Cell<Cell_Type> BArray< Cell_Type, Data_Type >::default_val ( ) const
```

#### 8.1.3.8 flush\_data()

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

#### 8.1.3.9 get\_cell()

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

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

# 8.1.3.12 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 >

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

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

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

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

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

#### 8.1.3.18 is dense()

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

Definition at line 240 of file barray-bones.hpp.

#### 8.1.3.19 is\_empty()

# 8.1.3.20 ncol()

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

## 8.1.3.21 nnozero()

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

# 8.1.3.22 nrow()

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

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

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

#### 8.1.3.25 operator\*=()

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

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

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

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

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

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

# 8.1.3.32 operator/=()

# 8.1.3.33 operator=() [1/2]

Move assignment.

# 8.1.3.34 operator=() [2/2]

Assignment constructor.

#### 8.1.3.35 operator==()

#### 8.1.3.36 out\_of\_range()

# 8.1.3.37 print()

#### 8.1.3.38 reserve()

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

#### 8.1.3.39 resize()

# 8.1.3.40 rm\_cell()

# 8.1.3.41 row()

# 8.1.3.42 set\_data()

Set the data object.

#### **Parameters**

data_	
delete_ <i>←</i>	
data_	

# 8.1.3.43 swap\_cells()

# 8.1.3.44 swap\_cols()

# 8.1.3.45 swap\_rows()

# 8.1.3.46 toggle\_cell()

# 8.1.3.47 toggle\_lock()

#### 8.1.3.48 transpose()

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

# 8.1.3.49 zero\_col()

# 8.1.3.50 zero\_row()

#### 8.1.4 Friends And Related Function Documentation

# 8.1.4.1 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.

## 8.1.4.2 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.

#### 8.1.5 Member Data Documentation

#### 8.1.5.1 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 54 of file barray-bones.hpp.

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

• include/barry/barray-bones.hpp

# 8.2 BArrayCell< Cell\_Type, Data\_Type > Class Template Reference

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

## **Public Member Functions**

- BArrayCell (BArray < Cell\_Type, Data\_Type > \*Array\_, size\_t i\_, size\_t 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

# 8.2.1 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.

# 8.2.2 Constructor & Destructor Documentation

# 8.2.2.1 BArrayCell()

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

#### 8.2.2.2 ~BArrayCell()

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

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

#### 8.2.3 Member Function Documentation

#### 8.2.3.1 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.

#### 8.2.3.2 operator\*=()

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

#### 8.2.3.3 operator+=()

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

### 8.2.3.4 operator-=()

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

#### 8.2.3.5 operator/=()

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

#### 8.2.3.6 operator=()

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

# 8.2.3.7 operator==()

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

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

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

# 8.3 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\_, size\_t i\_, size\_t 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
- 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

# 8.3.1 Detailed Description

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

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

# 8.3.2 Constructor & Destructor Documentation

#### 8.3.2.1 BArrayCell const()

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

#### 8.3.2.2 ~BArrayCell\_const()

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

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

# 8.3.3 Member Function Documentation

# 8.3.3.1 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.

#### 8.3.3.2 operator"!=()

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

#### 8.3.3.3 operator<()

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

#### 8.3.3.4 operator<=()

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

#### 8.3.3.5 operator==()

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

#### 8.3.3.6 operator>()

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

#### 8.3.3.7 operator>=()

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

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

- include/barry/barray-bones.hpp
- include/barry/barraycell-bones.hpp
- include/barry/barraycell-meat.hpp
- · include/barry/barrayrow-meat.hpp

# 8.4 BArrayDense< Cell\_Type, Data\_Type > Class Template Reference

Baseline class for binary arrays.

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

#### **Public Member Functions**

- bool operator== (const BArrayDense< Cell\_Type, Data\_Type > &Array\_)
- ∼BArrayDense ()
- void out\_of\_range (size\_t i, size\_t j) const
- Cell Type get cell (size ti, size tj, bool check bounds=true) const
- std::vector< Cell\_Type > get\_col\_vec (size\_t i, bool check\_bounds=true) const
- std::vector< Cell\_Type > get\_row\_vec (size\_t i, bool check\_bounds=true) const
- void get\_col\_vec (std::vector< Cell\_Type > \*x, size\_t i, bool check\_bounds=true) const
- void get row vec (std::vector< Cell Type > \*x, size t i, bool check bounds=true) const
- BArrayDenseRow< Cell\_Type, Data\_Type > & row (size\_t i, bool check\_bounds=true)
- const BArrayDenseRow\_const < Cell\_Type, Data\_Type > row (size\_t i, bool check\_bounds=true) const
- BArrayDenseCol< Cell Type, Data Type > & col (size t j, bool check bounds=true)
- const BArrayDenseCol\_const< Cell\_Type, Data\_Type > col (size\_t j, bool check\_bounds=true) const
- Entries < Cell\_Type > get\_entries () const

#### Get the edgelist.

- void transpose ()
- void clear (bool hard=true)
- void resize (size\_t N\_, size\_t M\_)
- void reserve ()
- void print (const char \*fmt=nullptr,...) const
- bool is\_dense () const noexcept
- const std::vector< Cell\_Type > & get\_data () const
- · const Cell Type rowsum (size ti) const
- const Cell\_Type colsum (size\_t i) const

#### Constructors

#### **Parameters**

N_	Number of rows
M_	Number of columns
source	An unsigned vector ranging from 0 to N_
target	An size_t vector ranging from 0 to M_
target	When true tries to add repeated observations.
value	Cell_Type defaul fill-in value (zero, by default.)

• BArrayDense ()

Zero-size array.

- BArrayDense (size\_t N\_, size\_t M\_, Cell\_Type value=static\_cast< Cell\_Type >(0))
   Empty array.
- BArrayDense (size\_t N\_, size\_t M\_, const std::vector< size\_t > &source, const std::vector< size\_t > &target, const std::vector< Cell\_Type > &value, bool add=true)

Edgelist with data.

BArrayDense (size\_t N\_, size\_t M\_, const std::vector< size\_t > &source, const std::vector< size\_t > &target, bool add=true)

Edgelist with no data (simpler)

- BArrayDense (const BArrayDense < Cell\_Type, Data\_Type > &Array\_, bool copy\_data=false)
   Copy constructor.
- BArrayDense< Cell\_Type, Data\_Type > & operator= (const BArrayDense< Cell\_Type, Data\_Type > &Array )

Assignment constructor.

 $\bullet \ \ \mathsf{BArrayDense} < \mathsf{Cell\_Type}, \ \mathsf{Data\_Type} > \&x) \ \mathsf{noexcept} \\$ 

Move operator

BArrayDense< Cell\_Type, Data\_Type > & operator= (BArrayDense< Cell\_Type, Data\_Type > &&x)
 noexcept

Move assignment.

void set\_data (Data\_Type \*data\_, bool delete\_data\_=false)

Set the data object.

- Data\_Type \* D\_ptr ()
- const Data\_Type \* D\_ptr () const
- Data\_Type & D ()
- const Data\_Type & D () const

# 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 (size\_t i, size\_t j, bool check\_bounds=true) const
- size\_t nrow () const noexcept
- size t ncol () const noexcept
- size\_t nnozero () const noexcept

Cell
 Cell\_Type > default\_val () 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 Wither check if the cell exists (before trying to delete/add), or, in the case of		
	swap_cells, check if either of both cells exists/don't exist.	

- BArrayDense< Cell\_Type, Data\_Type > & operator+= (const std::pair< size\_t, size\_t > &coords)
- BArrayDense< Cell Type, Data Type > & operator = (const std::pair < size t, size t > &coords)
- BArrayDenseCell< Cell\_Type, Data\_Type > operator() (size\_t i, size\_t j, bool check\_bounds=true)
- const Cell\_Type operator() (size\_t i, size\_t j, bool check\_bounds=true) const
- void rm cell (size t i, size t j, bool check bounds=true, bool check exists=true)
- void insert\_cell (size\_t i, size\_t j, const Cell< Cell\_Type > &v, bool check\_bounds, bool check\_exists)
- void insert cell (size ti, size ti, Cell Type v, bool check bounds, bool check exists)
- void swap\_cells (size\_t i0, size\_t j0, size\_t i1, size\_t j1, bool check\_bounds=true, int check\_exists=CHECK::BOTH, int \*report=nullptr)
- void toggle cell (size t i, size t i, bool check bounds=true, int check exists=EXISTS::UKNOWN)
- void toggle lock (size ti, size ti, bool check bounds=true)

### Column/row wise interchange

- void swap rows (size t i0, size t i1, bool check bounds=true)
- void swap cols (size t i0, size t i1, bool check bounds=true)
- void zero\_row (size\_t i, bool check\_bounds=true)
- void zero col (size t j, bool check bounds=true)

# **Arithmetic operators**

- BArrayDense< Cell\_Type, Data\_Type > & operator+= (const BArrayDense< Cell\_Type, Data\_Type > &rhs)
- BArrayDense< Cell\_Type, Data\_Type > & operator+= (const Cell\_Type &rhs)
- BArrayDense< Cell\_Type, Data\_Type > & operator== (const\_BArrayDense< Cell\_Type, Data\_Type > &rhs)
- BArrayDense< Cell\_Type, Data\_Type > & operator-= (const Cell\_Type &rhs)
- BArrayDense< Cell\_Type, Data\_Type > & operator/= (const Cell\_Type &rhs)
- BArrayDense < Cell\_Type, Data\_Type > & operator\*= (const Cell\_Type &rhs)

# **Public Attributes**

• bool visited = false

### **Friends**

- class BArrayDenseCell
   Cell\_Type, Data\_Type
- class BArrayDenseCol< Cell\_Type, Data\_Type >
- class BArrayDenseCol\_const< Cell\_Type, Data\_Type >
- class BArrayDenseRow
   Cell\_Type, Data\_Type
- class BArrayDenseRow\_const< Cell\_Type, Data\_Type >

# 8.4.1 Detailed Description

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

Baseline class for binary arrays.

BArrayDense class objects are arbitrary dense-arrays. The data is stored internally in the el member, which can be accessed using the member function get\_data(), by column.

#### **Template Parameters**

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

Definition at line 33 of file barraydense-bones.hpp.

# 8.4.2 Constructor & Destructor Documentation

# 8.4.2.1 BArrayDense() [1/6]

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

Zero-size array.

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

# 8.4.2.2 BArrayDense() [2/6]

Empty array.

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

#### 8.4.2.3 BArrayDense() [3/6]

Edgelist with data.

#### 8.4.2.4 BArrayDense() [4/6]

Edgelist with no data (simpler)

# 8.4.2.5 BArrayDense() [5/6]

Copy constructor.

# 8.4.2.6 BArrayDense() [6/6]

Move operator.

### 8.4.2.7 ~BArrayDense()

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

#### 8.4.3 Member Function Documentation

#### 8.4.3.1 clear()

# 8.4.3.2 col() [1/2]

Definition at line 490 of file barraydense-meat.hpp.

#### 8.4.3.3 col() [2/2]

Definition at line 476 of file barraydense-meat.hpp.

### 8.4.3.4 colsum()

# 8.4.3.5 D() [1/2]

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Data_Type& BArrayDense< Cell_Type, Data_Type >::D ( )
```

# 8.4.3.6 D() [2/2]

```
template<typename Cell_Type = bool, typename Data_Type = bool>
const Data_Type& BArrayDense< Cell_Type, Data_Type >::D ( ) const
```

# 8.4.3.7 D\_ptr() [1/2]

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Data_Type* BArrayDense< Cell_Type, Data_Type >::D_ptr ( )
```

# 8.4.3.8 D\_ptr() [2/2]

```
template<typename Cell_Type = bool, typename Data_Type = bool>
const Data_Type* BArrayDense< Cell_Type, Data_Type >::D_ptr ( ) const
```

# 8.4.3.9 default\_val()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Cell<Cell_Type> BArrayDense< Cell_Type, Data_Type >::default_val ( ) const
```

# 8.4.3.10 get\_cell()

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

#### 8.4.3.12 get col vec() [2/2]

# 8.4.3.13 get\_data()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
const std::vector< Cell_Type >& BArrayDense< Cell_Type, Data_Type >::get_data ( ) const
```

#### 8.4.3.14 get\_entries()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Entries<Cell_Type> BArrayDense< 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 >

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

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

# 8.4.3.17 insert\_cell() [1/2]

#### 8.4.3.18 insert cell() [2/2]

# 8.4.3.19 is\_dense()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
bool BArrayDense< Cell_Type, Data_Type >::is_dense ( ) const [inline], [noexcept]
```

Definition at line 256 of file barraydense-bones.hpp.

### 8.4.3.20 is\_empty()

# 8.4.3.21 ncol()

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

#### 8.4.3.22 nnozero()

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

#### 8.4.3.23 nrow()

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

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

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

#### 8.4.3.26 operator\*=()

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

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

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

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

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

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

#### 8.4.3.33 operator/=()

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

Move assignment.

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

Assignment constructor.

#### 8.4.3.36 operator==()

# 8.4.3.37 out\_of\_range()

# 8.4.3.38 print()

### 8.4.3.39 reserve()

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

#### 8.4.3.40 resize()

# 8.4.3.41 rm\_cell()

# 8.4.3.42 row() [1/2]

# 8.4.3.43 row() [2/2]

# 8.4.3.44 rowsum()

#### 8.4.3.45 set\_data()

Set the data object.

#### **Parameters**

data_	
delete_ <i>←</i>	
data_	

# 8.4.3.46 swap\_cells()

#### 8.4.3.47 swap\_cols()

# 8.4.3.48 swap\_rows()

# 8.4.3.49 toggle\_cell()

# 8.4.3.50 toggle\_lock()

#### 8.4.3.51 transpose()

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

#### 8.4.3.52 zero col()

# 8.4.3.53 zero\_row()

# 8.4.4 Friends And Related Function Documentation

#### 8.4.4.1 BArrayDenseCell< Cell\_Type, Data\_Type >

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

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

# 8.4.4.2 BArrayDenseCol< Cell\_Type, Data\_Type>

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

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

## 8.4.4.3 BArrayDenseCol\_const< Cell\_Type, Data\_Type >

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

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

# 8.4.4.4 BArrayDenseRow < Cell\_Type, Data\_Type >

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

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

#### 8.4.4.5 BArrayDenseRow\_const< Cell\_Type, Data\_Type >

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

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

#### 8.4.5 Member Data Documentation

#### 8.4.5.1 visited

```
template<typename Cell_Type = bool, typename Data_Type = bool>
bool BArrayDense< 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 63 of file barraydense-bones.hpp.

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

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

# 8.5 BArrayDenseCell< Cell\_Type, Data\_Type > Class Template Reference

#include <barraydensecell-bones.hpp>

#### **Public Member Functions**

- $\bullet \ \, \mathsf{BArrayDenseCell} \ (\mathsf{BArrayDense} < \mathsf{Cell\_Type}, \mathsf{Data\_Type} > *\mathsf{Array\_}, \mathsf{size\_t} \ i\_, \mathsf{size\_t} \ j\_, \mathsf{bool} \ \mathsf{check\_bounds} = \mathsf{true})$
- BArrayDenseCell< Cell\_Type, Data\_Type > & operator= (const BArrayDenseCell< Cell\_Type, Data\_Type > &other)
- ∼BArrayDenseCell ()
- 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

#### **Friends**

- class BArrayDense< Cell\_Type, Data\_Type >
- class BArrayDenseCol < Cell\_Type, Data\_Type >
- class BArrayDenseCol\_const< Cell\_Type, Data\_Type >

# 8.5.1 Detailed Description

template<typename Cell\_Type = bool, typename Data\_Type = bool> class BArrayDenseCell< Cell\_Type, Data\_Type >

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

# 8.5.2 Constructor & Destructor Documentation

# 8.5.2.1 BArrayDenseCell()

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

#### 8.5.2.2 ~BArrayDenseCell()

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

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

#### 8.5.3 Member Function Documentation

#### 8.5.3.1 operator Cell\_Type()

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

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

#### 8.5.3.2 operator\*=()

Definition at line 52 of file barraydensecell-meat.hpp.

#### 8.5.3.3 operator+=()

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

#### 8.5.3.4 operator-=()

Definition at line 43 of file barraydensecell-meat.hpp.

# 8.5.3.5 operator/=()

Definition at line 62 of file barraydensecell-meat.hpp.

# 8.5.3.6 operator=() [1/2]

Definition at line 9 of file barraydensecell-meat.hpp.

# 8.5.3.7 operator=() [2/2]

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

#### 8.5.3.8 operator==()

Definition at line 77 of file barraydensecell-meat.hpp.

#### 8.5.4 Friends And Related Function Documentation

# 8.5.4.1 BArrayDense < Cell\_Type, Data\_Type >

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

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

# 8.5.4.2 BArrayDenseCol < Cell\_Type, Data\_Type >

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

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

# 8.5.4.3 BArrayDenseCol\_const< Cell\_Type, Data\_Type >

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

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

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

- include/barry/barraydense-bones.hpp
- include/barry/barraydensecell-bones.hpp
- include/barry/barraydensecell-meat.hpp

# 8.6 BArrayDenseCell\_const< Cell\_Type, Data\_Type > Class Template Reference

# 8.6.1 Detailed Description

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

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

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

• include/barry/barraydense-bones.hpp

# 8.7 BArrayDenseCol< Cell\_Type, Data\_Type > Class Template Reference

#include <barraydensecol-bones.hpp>

#### **Public Member Functions**

- BArrayDenseCol (BArrayDense < Cell\_Type, Data\_Type > &array\_, size\_t j)
- Col\_type< Cell\_Type >::iterator & begin ()
- Col\_type< Cell\_Type >::iterator & end ()
- size\_t size () const noexcept
- std::pair< size\_t, Cell\_Type \* > & operator() (size\_t i)

# **Friends**

- class BArrayDense< Cell\_Type, Data\_Type >
- class BArrayDenseCell
   Cell Type, Data Type
- $\bullet \ \ {\it class BArrayDenseCell\_const} < {\it Cell\_Type}, \ {\it Data\_Type} >$

# 8.7.1 Detailed Description

```
\label{template} \mbox{typename Cell_Type = bool, typename Data_Type = bool} > \mbox{class BArrayDenseCol} < \mbox{Cell_Type, Data_Type} > \mbox{}
```

Definition at line 9 of file barraydensecol-bones.hpp.

## 8.7.2 Constructor & Destructor Documentation

# 8.7.2.1 BArrayDenseCol()

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

#### 8.7.3 Member Function Documentation

# 8.7.3.1 begin()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Col_type<Cell_Type>::iterator& BArrayDenseCol< Cell_Type, Data_Type >::begin ( ) [inline]
```

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

#### 8.7.3.2 end()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Col_type<Cell_Type>::iterator& BArrayDenseCol< Cell_Type, Data_Type >::end ( ) [inline]
```

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

# 8.7.3.3 operator()()

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

# 8.7.3.4 size()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
size_t BArrayDenseCol< Cell_Type, Data_Type >::size ( ) const [inline], [noexcept]
```

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

# 8.7.4 Friends And Related Function Documentation

# 8.7.4.1 BArrayDense < Cell\_Type, Data\_Type >

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

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

# 8.7.4.2 BArrayDenseCell< Cell\_Type, Data\_Type >

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

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

# 8.7.4.3 BArrayDenseCell\_const< Cell\_Type, Data\_Type >

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

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

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

- include/barry/barraydense-bones.hpp
- include/barry/barraydensecol-bones.hpp

# 8.8 BArrayDenseCol\_const< Cell\_Type, Data\_Type > Class Template Reference

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

#### **Public Member Functions**

- BArrayDenseCol\_const (const BArrayDense< Cell\_Type, Data\_Type > &array\_, size\_t j)
- Col\_type< Cell\_Type >::iterator begin ()
- Col\_type< Cell\_Type >::iterator end ()
- size t size () const noexcept
- const std::pair< size\_t, Cell\_Type \* > operator() (size\_t i) const

## **Friends**

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

# 8.8.1 Detailed Description

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

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

# 8.8.2 Constructor & Destructor Documentation

## 8.8.2.1 BArrayDenseCol\_const()

Definition at line 80 of file barraydensecol-bones.hpp.

# 8.8.3 Member Function Documentation

# 8.8.3.1 begin()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Col_type<Cell_Type>::iterator BArrayDenseCol_const< Cell_Type, Data_Type >::begin ( ) [inline]
```

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

# 8.8.3.2 end()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Col_type<Cell_Type>::iterator BArrayDenseCol_const< Cell_Type, Data_Type >::end ( ) [inline]
```

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

## 8.8.3.3 operator()()

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

#### 8.8.3.4 size()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
size_t BArrayDenseCol_const< Cell_Type, Data_Type >::size ( ) const [inline], [noexcept]
```

Definition at line 107 of file barraydensecol-bones.hpp.

#### 8.8.4 Friends And Related Function Documentation

## 8.8.4.1 BArrayDenseCell< Cell\_Type, Data\_Type >

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

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

# 8.8.4.2 BArrayDenseCell\_const < Cell\_Type, Data\_Type >

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

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

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

- include/barry/barraydense-bones.hpp
- include/barry/barraydensecol-bones.hpp

# 8.9 BArrayDenseRow< Cell\_Type, Data\_Type > Class Template Reference

#include <barraydenserow-bones.hpp>

# **Public Member Functions**

- BArrayDenseRow (BArrayDense < Cell\_Type, Data\_Type > &array\_, size\_t i)
- Row\_type< Cell\_Type >::iterator & begin ()
- Row\_type< Cell\_Type >::iterator & end ()
- size\_t size () const noexcept
- std::pair< size\_t, Cell< Cell\_Type > > & operator() (size\_t i)

#### Friends

- class BArrayDense< Cell\_Type, Data\_Type >
- class BArrayDenseCell
   Cell\_Type, Data\_Type
- class BArrayDenseCell\_const< Cell\_Type, Data\_Type >

# 8.9.1 Detailed Description

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

Definition at line 9 of file barraydenserow-bones.hpp.

#### 8.9.2 Constructor & Destructor Documentation

## 8.9.2.1 BArrayDenseRow()

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

# 8.9.3 Member Function Documentation

# 8.9.3.1 begin()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Row_type<Cell_Type>::iterator& BArrayDenseRow< Cell_Type, Data_Type >::begin ( ) [inline]
```

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

# 8.9.3.2 end()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Row_type<Cell_Type>::iterator& BArrayDenseRow< Cell_Type, Data_Type >::end () [inline]
```

Definition at line 53 of file barraydenserow-bones.hpp.

## 8.9.3.3 operator()()

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

## 8.9.3.4 size()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
size_t BArrayDenseRow< Cell_Type, Data_Type >::size ( ) const [inline], [noexcept]
```

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

# 8.9.4 Friends And Related Function Documentation

# 8.9.4.1 BArrayDense< Cell\_Type, Data\_Type>

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

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

# 8.9.4.2 BArrayDenseCell< Cell\_Type, Data\_Type>

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

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

## 8.9.4.3 BArrayDenseCell\_const< Cell\_Type, Data\_Type >

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

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

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

- include/barry/barraydense-bones.hpp
- include/barry/barraydenserow-bones.hpp

# 8.10 BArrayDenseRow\_const< Cell\_Type, Data\_Type > Class Template Reference

#include <barraydenserow-bones.hpp>

## **Public Member Functions**

- BArrayDenseRow\_const (const BArrayDense< Cell\_Type, Data\_Type > &array\_, size\_t i)
- Row\_type< Cell\_Type >::const\_iterator begin () const
- Row\_type< Cell\_Type >::const\_iterator end () const
- size\_t size () const noexcept
- const std::pair< size\_t, Cell< Cell\_Type > > operator() (size\_t i) const

## **Friends**

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

# 8.10.1 Detailed Description

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

Definition at line 80 of file barraydenserow-bones.hpp.

# 8.10.2 Constructor & Destructor Documentation

# 8.10.2.1 BArrayDenseRow\_const()

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

# 8.10.3 Member Function Documentation

## 8.10.3.1 begin()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Row_type< Cell_Type >::const_iterator BArrayDenseRow_const< Cell_Type, Data_Type >::begin ( )
const [inline]
```

Definition at line 108 of file barraydenserow-bones.hpp.

# 8.10.3.2 end()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Row_type< Cell_Type >::const_iterator BArrayDenseRow_const< Cell_Type, Data_Type >::end ( )
const [inline]
```

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

## 8.10.3.3 operator()()

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

## 8.10.3.4 size()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
size_t BArrayDenseRow_const< Cell_Type, Data_Type >::size ( ) const [inline], [noexcept]
```

Definition at line 118 of file barraydenserow-bones.hpp.

# 8.10.4 Friends And Related Function Documentation

# 8.10.4.1 BArrayDenseCell< Cell\_Type, Data\_Type>

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

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

## 8.10.4.2 BArrayDenseCell\_const< Cell\_Type, Data\_Type >

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

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

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

- · include/barry/barraydense-bones.hpp
- include/barry/barraydenserow-bones.hpp

# 8.11 BArrayRow < Cell\_Type, Data\_Type > Class Template Reference

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

## **Public Member Functions**

- BArrayRow (BArray < Cell\_Type, Data\_Type > \*Array\_, size\_t i\_, bool check\_bounds=true)
- ∼BArrayRow ()
- void operator= (const BArrayRow< Cell\_Type, Data\_Type > &val)
- void operator+= (const BArrayRow< Cell\_Type, Data\_Type > &val)
- void operator== (const BArrayRow< Cell\_Type, Data\_Type > &val)
- $\bullet \ \ \mathsf{void} \ \mathsf{operator} *= (\mathsf{const} \ \mathsf{BArrayRow} < \mathsf{Cell\_Type}, \ \mathsf{Data\_Type} > \mathsf{\&val})$
- void operator/= (const BArrayRow< Cell\_Type, Data\_Type > &val)
- operator BArrayRow< Cell\_Type, Data\_Type > () const
- bool operator== (const BArrayRow< Cell\_Type, Data\_Type > &val) const

# 8.11.1 Detailed Description

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

Definition at line 5 of file barrayrow-bones.hpp.

# 8.11.2 Constructor & Destructor Documentation

# 8.11.2.1 BArrayRow()

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

## 8.11.2.2 ∼BArrayRow()

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

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

# 8.11.3 Member Function Documentation

# 8.11.3.1 operator BArrayRow< Cell\_Type, Data\_Type >()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
BArrayRow< Cell_Type, Data_Type >::operator BArrayRow< Cell_Type, Data_Type > ( ) const
```

# 8.11.3.2 operator\*=()

# 8.11.3.3 operator+=()

#### 8.11.3.4 operator-=()

## 8.11.3.5 operator/=()

## 8.11.3.6 operator=()

# 8.11.3.7 operator==()

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

• include/barry/barrayrow-bones.hpp

# 8.12 BArrayRow\_const< Cell\_Type, Data\_Type > Class Template Reference

#include <barrayrow-bones.hpp>

# **Public Member Functions**

- BArrayRow\_const (const BArray < Cell\_Type, Data\_Type > \*Array\_, size\_t i\_, bool check\_bounds=true)
- ∼BArrayRow\_const ()
- operator BArrayRow\_const< Cell\_Type, Data\_Type > () const
- bool operator== (const BArrayRow\_const< Cell\_Type, Data\_Type > &val) const
- bool operator!= (const BArrayRow const< Cell Type, Data Type > &val) const
- bool operator< (const BArrayRow\_const< Cell\_Type, Data\_Type > &val) const
- bool operator> (const BArrayRow\_const< Cell\_Type, Data\_Type > &val) const
- bool operator<= (const BArrayRow\_const< Cell\_Type, Data\_Type > &val) const
- bool operator>= (const BArrayRow\_const< Cell\_Type, Data\_Type > &val) const

# 8.12.1 Detailed Description

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

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

# 8.12.2 Constructor & Destructor Documentation

## 8.12.2.1 BArrayRow\_const()

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

## 8.12.2.2 ~BArrayRow\_const()

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

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

## 8.12.3 Member Function Documentation

# 8.12.3.1 operator BArrayRow\_const< Cell\_Type, Data\_Type >()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
BArrayRow_const< Cell_Type, Data_Type >::operator BArrayRow_const< Cell_Type, Data_Type > ( )
const
```

# 8.12.3.2 operator"!=()

## 8.12.3.3 operator<()

#### 8.12.3.4 operator<=()

#### 8.12.3.5 operator==()

# 8.12.3.6 operator>()

# 8.12.3.7 operator>=()

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

• include/barry/barrayrow-bones.hpp

# 8.13 BArrayVector< Cell Type, Data Type > Class Template Reference

Row or column of a BArray

#include <barrayvector-bones.hpp>

# **Public Member Functions**

- BArrayVector (BArray < Cell\_Type, Data\_Type > \*Array\_, size\_t &dim\_ size\_t &i\_, bool check\_bounds=true)

  Construct a new BArrayVector object.
- ∼BArrayVector ()
- · bool is row () const noexcept
- bool is\_col () const noexcept
- size\_t size () const noexcept
- std::vector< Cell\_Type >::const\_iterator begin () noexcept
- std::vector< Cell\_Type >::const\_iterator end () noexcept
- 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 std::vector< Cell\_Type > () const
- bool operator== (const Cell\_Type &val) const

# 8.13.1 Detailed Description

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

Row or column of a BArray

**Template Parameters** 

Cell_Type	
Data_Type	

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

# 8.13.2 Constructor & Destructor Documentation

# 8.13.2.1 BArrayVector()

Construct a new BArrayVector object.

# **Parameters**

Array_	Pointer to a BArray object
dim_	Dimension. 0 means row and 1 means column.
i_	Element to point.
check_bounds	When true, check boundaries.

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

## 8.13.2.2 ~BArrayVector()

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

Definition at line 53 of file barrayvector-bones.hpp.

## 8.13.3 Member Function Documentation

# 8.13.3.1 begin()

```
template<typename Cell_Type , typename Data_Type >
std::vector< Cell_Type >::const_iterator BArrayVector< Cell_Type, Data_Type >::begin [inline],
[noexcept]
```

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

## 8.13.3.2 end()

```
template<typename Cell_Type , typename Data_Type >
std::vector< Cell_Type >::const_iterator BArrayVector< Cell_Type, Data_Type >::end [inline],
[noexcept]
```

Definition at line 64 of file barrayvector-meat.hpp.

# 8.13.3.3 is\_col()

```
template<typename Cell_Type , typename Data_Type >
bool BArrayVector< Cell_Type, Data_Type >::is_col [inline], [noexcept]
```

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

# 8.13.3.4 is\_row()

```
template<typename Cell_Type , typename Data_Type >
bool BArrayVector< Cell_Type, Data_Type >::is_row [inline], [noexcept]
```

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

# 8.13.3.5 operator std::vector< Cell\_Type >()

```
template<typename Cell_Type , typename Data_Type >
BArrayVector< Cell_Type, Data_Type >::operator std::vector< Cell_Type > [inline]
```

Definition at line 175 of file barrayvector-meat.hpp.

# 8.13.3.6 operator \*= ()

Definition at line 133 of file barrayvector-meat.hpp.

## 8.13.3.7 operator+=()

Definition at line 91 of file barrayvector-meat.hpp.

## 8.13.3.8 operator-=()

Definition at line 112 of file barrayvector-meat.hpp.

## 8.13.3.9 operator/=()

Definition at line 154 of file barrayvector-meat.hpp.

# 8.13.3.10 operator=()

Definition at line 69 of file barrayvector-meat.hpp.

# 8.13.3.11 operator==()

Definition at line 185 of file barrayvector-meat.hpp.

## 8.13.3.12 size()

```
template<typename Cell_Type , typename Data_Type >
size_t BArrayVector< Cell_Type, Data_Type >::size [inline], [noexcept]
```

Definition at line 39 of file barrayvector-meat.hpp.

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

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

# 8.14 BArrayVector\_const< Cell\_Type, Data\_Type > Class Template Reference

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

## **Public Member Functions**

- BArrayVector\_const (const BArray< Cell\_Type, Data\_Type > \*Array\_, size\_t &dim\_ size\_t &i\_, bool check\_bounds=true)
- ∼BArrayVector\_const ()
- · bool is\_row () const noexcept
- bool is\_col () const noexcept
- size\_t size () const noexcept
- std::vector< Cell Type >::const iterator begin () noexcept
- std::vector< Cell Type >::const iterator end () noexcept
- operator std::vector< Cell Type > () const
- bool operator== (const Cell\_Type &val) const
- bool operator!= (const Cell\_Type &val) const
- bool operator< (const Cell\_Type &val) const
- bool operator> (const Cell\_Type &val) const
- bool operator<= (const Cell\_Type &val) const
- bool operator>= (const Cell Type &val) const

# 8.14.1 Detailed Description

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

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

## 8.14.2 Constructor & Destructor Documentation

# 8.14.2.1 BArrayVector\_const()

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

#### 8.14.2.2 ~BArrayVector\_const()

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

Definition at line 108 of file barrayvector-bones.hpp.

# 8.14.3 Member Function Documentation

# 8.14.3.1 begin()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
std::vector< Cell_Type >::const_iterator BArrayVector_const< Cell_Type, Data_Type >::begin (
) [noexcept]
```

# 8.14.3.2 end()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
std::vector< Cell_Type >::const_iterator BArrayVector_const< Cell_Type, Data_Type >::end ( )
[noexcept]
```

# 8.14.3.3 is\_col()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
bool BArrayVector_const< Cell_Type, Data_Type >::is_col () const [noexcept]
```

## 8.14.3.4 is\_row()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
bool BArrayVector_const< Cell_Type, Data_Type >::is_row ( ) const [noexcept]
```

## 8.14.3.5 operator std::vector< Cell\_Type >()

```
template<typename Cell_Type , typename Data_Type >
BArrayVector_const< Cell_Type, Data_Type >::operator std::vector< Cell_Type > [inline]
```

Definition at line 212 of file barrayvector-meat.hpp.

## 8.14.3.6 operator"!=()

Definition at line 249 of file barrayvector-meat.hpp.

# 8.14.3.7 operator<()

Definition at line 254 of file barrayvector-meat.hpp.

# 8.14.3.8 operator<=()

Definition at line 281 of file barrayvector-meat.hpp.

# 8.14.3.9 operator==()

Definition at line 222 of file barrayvector-meat.hpp.

# 8.14.3.10 operator>()

Definition at line 308 of file barrayvector-meat.hpp.

## 8.14.3.11 operator>=()

Definition at line 315 of file barrayvector-meat.hpp.

# 8.14.3.12 size()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
size_t BArrayVector_const< Cell_Type, Data_Type >::size ( ) const [noexcept]
```

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

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

# 8.15 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, bool active\_=true)
- ∼Cell ()
- Cell (const Cell
   Cell\_Type > &arg)
- Cell< Cell\_Type > & operator= (const 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
- bool operator== (const Cell< Cell\_Type > &rhs) const
- bool operator!= (const Cell< Cell\_Type > &rhs) const
- void add (double x)
- void add (size\_t x)
- void add (int x)
- Cell ()
- Cell ()
- Cell ()

# **Public Attributes**

- Cell\_Type value
- bool visited
- · bool active

# 8.15.1 Detailed Description

```
\label{eq:class} \begin{tabular}{ll} template < class Cell\_Type > \\ class Cell < Cell\_Type > \\ \end{tabular}
```

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

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

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

## 8.15.2 Constructor & Destructor Documentation

# 8.15.2.1 Cell() [1/7]

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

# 8.15.2.2 Cell() [2/7]

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

# 8.15.2.3 $\sim$ Cell()

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

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

# 8.15.2.4 Cell() [3/7]

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

# 8.15.2.5 Cell() [4/7]

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

# 8.15.2.6 Cell() [5/7]

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

Definition at line 62 of file cell-meat.hpp.

# 8.15.2.7 Cell() [6/7]

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

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

# 8.15.2.8 Cell() [7/7]

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

Definition at line 64 of file cell-meat.hpp.

# 8.15.3 Member Function Documentation

# 8.15.3.1 add() [1/4]

# 8.15.3.2 add() [2/4]

Definition at line 42 of file cell-meat.hpp.

# 8.15.3.3 add() [3/4]

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

Definition at line 52 of file cell-meat.hpp.

# 8.15.3.4 add() [4/4]

Definition at line 47 of file cell-meat.hpp.

# 8.15.3.5 operator Cell\_Type()

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

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

# 8.15.3.6 operator"!=()

Definition at line 31 of file cell-meat.hpp.

# 8.15.3.7 operator=() [1/2]

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

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

Definition at line 5 of file cell-meat.hpp.

# 8.15.3.9 operator==()

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

# 8.15.4 Member Data Documentation

# 8.15.4.1 active

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

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

# 8.15.4.2 value

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

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

## 8.15.4.3 visited

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

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

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

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

# 8.16 Cell\_const< Cell\_Type > Class Template Reference

# 8.16.1 Detailed Description

```
\label{lem:const} \begin{tabular}{ll} template < typename Cell_Type > \\ class Cell_const < Cell_Type > \\ \end{tabular}
```

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

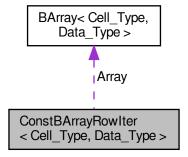
The documentation for this class was generated from the following file:

• include/barry/barray-meat.hpp

# 8.17 ConstBArrayRowlter< Cell\_Type, Data\_Type > Class Template Reference

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

Collaboration diagram for ConstBArrayRowlter< Cell\_Type, Data\_Type >:



# **Public Member Functions**

- ConstBArrayRowlter (const BArray< Cell\_Type, Data\_Type > \*Array\_)
- ∼ConstBArrayRowIter ()

## **Public Attributes**

- · size t current row
- size\_t current\_col
- Row\_type< Cell\_Type >::const\_iterator iter
- const BArray
   Cell\_Type, Data\_Type > \* Array

# 8.17.1 Detailed Description

```
\label{template} \begin{tabular}{ll} template < typename Cell_Type, typename Data_Type > \\ class ConstBArrayRowlter < Cell_Type, Data_Type > \\ \end{tabular}
```

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

# 8.17.2 Constructor & Destructor Documentation

# 8.17.2.1 ConstBArrayRowIter()

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

## 8.17.2.2 ∼ConstBArrayRowlter()

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

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

## 8.17.3 Member Data Documentation

## 8.17.3.1 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.

## 8.17.3.2 current\_col

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

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

## 8.17.3.3 current\_row

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

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

#### 8.17.3.4 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

# 8.18 Counter< Array\_Type, Data\_Type > Class Template Reference

A counter function based on change statistics.

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

# **Public Member Functions**

- ∼Counter ()
- double count (Array\_Type &Array, size\_t i, size\_t j)
- double init (Array\_Type &Array, size\_t i, size\_t j)
- std::string get\_name () const
- std::string get\_description () const

Creator passing a counter and an initializer

#### **Parameters**

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

- Counter ()
- Counter\_fun\_type
   Array\_Type, Data\_Type > count\_fun\_, Counter\_fun\_type
   Array\_Type, Data\_Type > init\_fun\_, Hasher\_fun\_type
   Array\_Type, Data\_Type > hasher\_fun\_, Data\_Type data\_, std::string name\_="", std::string desc\_="")
- Counter (const Counter < Array\_Type, Data\_Type > &counter\_)
   Copy constructor.
- Counter (Counter < Array\_Type, Data\_Type > &&counter\_) noexcept

Move constructor

- Counter< Array\_Type, Data\_Type > operator= (const Counter< Array\_Type, Data\_Type > &counter\_)
   Copy assignment.
- Counter< Array\_Type, Data\_Type > & operator= (Counter< Array\_Type, Data\_Type > &&counter\_)
   noexcept

Move assignment.

void set\_hasher (Hasher\_fun\_type< Array\_Type, Data\_Type > fun)

Get and set the hasher function.

Hasher\_fun\_type< Array\_Type, Data\_Type > get\_hasher ()

# **Public Attributes**

- Counter\_fun\_type
   Array\_Type, Data\_Type > count\_fun
- Counter\_fun\_type< Array\_Type, Data\_Type > init\_fun
- Hasher\_fun\_type
   Array\_Type, Data\_Type > hasher\_fun
- Data\_Type data
- std::string name = ""
- std::string desc = ""

# 8.18.1 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 35 of file counters-bones.hpp.

# 8.18.2 Constructor & Destructor Documentation

# 8.18.2.1 Counter() [1/4]

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

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

## 8.18.2.2 Counter() [2/4]

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

## 8.18.2.3 Counter() [3/4]

Copy constructor.

## 8.18.2.4 Counter() [4/4]

Move constructor.

## 8.18.2.5 ∼Counter()

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

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

## 8.18.3 Member Function Documentation

# 8.18.3.1 count()

# 8.18.3.2 get\_description()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
std::string Counter< Array_Type, Data_Type >::get_description ( ) const
```

## 8.18.3.3 get\_hasher()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Hasher_fun_type<Array_Type, Data_Type> Counter< Array_Type, Data_Type >::get_hasher ( )
```

## 8.18.3.4 get\_name()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
std::string Counter< Array_Type, Data_Type >::get_name ( ) const
```

# 8.18.3.5 init()

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

Copy assignment.

# 8.18.3.7 operator=() [2/2]

Move assignment.

# 8.18.3.8 set\_hasher()

Get and set the hasher function.

The hasher function is used to characterize the support of the array. This way, if possible, the support enumeration is recycled.

**Parameters** 

fun

## 8.18.4 Member Data Documentation

# 8.18.4.1 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 38 of file counters-bones.hpp.

#### 8.18.4.2 data

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

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

#### 8.18.4.3 desc

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

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

## 8.18.4.4 hasher\_fun

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Hasher_fun_type<Array_Type,Data_Type> Counter< Array_Type, Data_Type >::hasher_fun
```

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

# 8.18.4.5 init\_fun

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

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

# 8.18.4.6 name

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

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

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

• include/barry/counters-bones.hpp

# 8.19 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\_)

Copy constructor.

Counters (Counters < Array\_Type, Data\_Type > &&counters\_) noexcept

Move constructor.

Counters < Array\_Type, Data\_Type > operator= (const Counters < Array\_Type, Data\_Type > &counter\_)
 Copy assignment constructor.

Counters< Array\_Type, Data\_Type > & operator= (Counters< Array\_Type, Data\_Type > &&counter\_)
 noexcept

Move assignment constructor.

Counter< Array\_Type, Data\_Type > & operator[] (size\_t idx)

Returns a pointer to a particular counter.

• std::size\_t size () const noexcept

Number of counters in the set.

- void add\_counter (Counter< Array\_Type, Data\_Type > counter)
- std::vector< std::string > get\_names () const
- std::vector< std::string > get\_descriptions () const
- std::vector< double > gen\_hash (const Array\_Type &array, bool add\_dims=true)

Generates a hash for the given array according to the counters.

void add\_hash (Hasher\_fun\_type< Array\_Type, Data\_Type > fun\_)

# 8.19.1 Detailed Description

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

Vector of counters.

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

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

## 8.19.2 Constructor & Destructor Documentation

# 8.19.2.1 Counters() [1/3]

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

# 8.19.2.2 ~Counters()

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

Definition at line 120 of file counters-bones.hpp.

## 8.19.2.3 Counters() [2/3]

Copy constructor.

# Parameters



# 8.19.2.4 Counters() [3/3]

Move constructor.

#### **Parameters**

counters⇔

# 8.19.3 Member Function Documentation

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

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

## 8.19.3.3 add hash()

## 8.19.3.4 gen\_hash()

Generates a hash for the given array according to the counters.

## **Parameters**

array	
add_dims	When true (default) the dimmension of the array will be added to the hash.

# Returns

std::vector< double > That can be hashed later.

## 8.19.3.5 get\_descriptions()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
std::vector< std::string > Counters< Array_Type, Data_Type >::get_descriptions ( ) const
```

# 8.19.3.6 get\_names()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
std::vector< std::string > Counters< Array_Type, Data_Type >::get_names ( ) const
```

# 8.19.3.7 operator=() [1/2]

Copy assignment constructor.

## **Parameters**



#### Returns

Counters<Array\_Type,Data\_Type>

# 8.19.3.8 operator=() [2/2]

Move assignment constructor.

## **Parameters**



## Returns

```
Counters<Array_Type,Data_Type>&
```

# 8.19.3.9 operator[]()

Returns a pointer to a particular counter.

#### **Parameters**

```
idx Id of the counter
```

## Returns

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

# 8.19.3.10 size()

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

Number of counters in the set.

# Returns

size\_t

Definition at line 164 of file counters-bones.hpp.

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

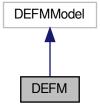
• include/barry/counters-bones.hpp

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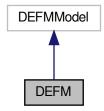
# 8.20 DEFM Class Reference

#include <defm-bones.hpp>

Inheritance diagram for DEFM:



Collaboration diagram for DEFM:



#### **Public Member Functions**

- DEFM (const int \*id, const int \*y, const double \*x, size\_t id\_length, size\_t y\_ncol, size\_t x\_ncol, size\_t m\_
   order)
- DEFMModel & get\_model ()
- void init ()
- double likelihood (std::vector< double > &par, bool as\_log=false)
- void simulate (std::vector< double > par, int \*y\_out)
- size\_t get\_n\_y () const
- size\_t get\_n\_obs () const
- size\_t get\_n\_covars () const
- size\_t get\_m\_order () const
- size\_t get\_n\_rows () const
- const int \* get\_Y () const
- const int \* get\_ID () const
- const double \* get\_X () const

```
barry::FreqTable < int > motif_census (std::vector < size_t > idx)
std::vector < double > logodds (const std::vector < double > &par, size_t i, size_t j)
void set_names (std::vector < std::string > Y_names_, std::vector < std::string > X_names_)
const std::vector < std::string > & get_Y_names () const
const std::vector < std::string > & get_X_names () const
void print () const
std::vector < bool > is_motif ()
```

# 8.20.1 Detailed Description

Definition at line 4 of file defm-bones.hpp.

#### 8.20.2 Constructor & Destructor Documentation

#### 8.20.2.1 DEFM()

Definition at line 104 of file defm-meat.hpp.

## 8.20.3 Member Function Documentation

```
8.20.3.1 get_ID()
```

```
const int * DEFM::get_ID ( ) const [inline]
```

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

# 8.20.3.2 get\_m\_order()

```
size_t DEFM::get_m_order ( ) const [inline]
```

Definition at line 244 of file defm-meat.hpp.

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## 8.20.3.3 get\_model()

```
DEFMModel& DEFM::get_model ( ) [inline]
```

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

#### 8.20.3.4 get\_n\_covars()

```
size_t DEFM::get_n_covars ( ) const [inline]
```

Definition at line 239 of file defm-meat.hpp.

#### 8.20.3.5 get\_n\_obs()

```
size_t DEFM::get_n_obs ( ) const [inline]
```

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

# 8.20.3.6 get\_n\_rows()

```
size_t DEFM::get_n_rows ( ) const [inline]
```

Definition at line 249 of file defm-meat.hpp.

# 8.20.3.7 get\_n\_y()

```
size_t DEFM::get_n_y ( ) const [inline]
```

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

# 8.20.3.8 get\_X()

```
const double * DEFM::get_X ( ) const [inline]
```

Definition at line 264 of file defm-meat.hpp.

# 8.20.3.9 get\_X\_names()

```
const std::vector< std::string > & DEFM::get_X_names ( ) const [inline]
```

Definition at line 371 of file defm-meat.hpp.

#### 8.20.3.10 get\_Y()

```
const int * DEFM::get_Y ( ) const [inline]
```

Definition at line 254 of file defm-meat.hpp.

# 8.20.3.11 get\_Y\_names()

```
const std::vector< std::string > & DEFM::get_Y_names ( ) const [inline]
```

Definition at line 367 of file defm-meat.hpp.

# 8.20.3.12 init()

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

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

# 8.20.3.13 is\_motif()

```
std::vector< bool > DEFM::is_motif ( ) [inline]
```

Definition at line 388 of file defm-meat.hpp.

# 8.20.3.14 likelihood()

```
double DEFM::likelihood (
          std::vector< double > & par,
          bool as_log = false )
```

8.20 DEFM Class Reference 141

#### 8.20.3.15 logodds()

Definition at line 308 of file defm-meat.hpp.

# 8.20.3.16 motif\_census()

```
barry::FreqTable< int > DEFM::motif_census (  std::vector < size_t > \mathit{idx} ) \quad [inline]
```

Definition at line 270 of file defm-meat.hpp.

#### 8.20.3.17 print()

```
void DEFM::print ( ) const [inline]
```

Definition at line 375 of file defm-meat.hpp.

## 8.20.3.18 set\_names()

```
void DEFM::set_names ( std::vector < std::string > \textit{Y}\_names\_, \\ std::vector < std::string > \textit{X}\_names\_ ) \quad [inline]
```

Definition at line 350 of file defm-meat.hpp.

## 8.20.3.19 simulate()

```
void DEFM::simulate (
          std::vector< double > par,
          int * y_out ) [inline]
```

Definition at line 38 of file defm-meat.hpp.

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

- include/barry/models/defm/defm-bones.hpp
- include/barry/models/defm/defm-meat.hpp

# 8.21 DEFMCounterData Class Reference

Data class used to store arbitrary size\_t or double vectors.

```
#include <defm-types.hpp>
```

#### **Public Member Functions**

- DEFMCounterData ()
- DEFMCounterData (const std::vector< size\_t > indices\_, const std::vector< double > numbers\_, const std::vector< bool > logical\_, bool is\_motif\_=true)
- size\_t idx (size\_t i) const
- double num (size\_t i) const
- · bool is true (size ti) const
- ∼DEFMCounterData ()

#### **Public Attributes**

- std::vector< size\_t > indices
- std::vector< double > numbers
- std::vector< bool > logical
- · bool is\_motif

If false, then is a logit intercept.

# 8.21.1 Detailed Description

Data class used to store arbitrary size\_t or double vectors.

Definition at line 64 of file defm-types.hpp.

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

• include/barry/models/defm/defm-types.hpp

# 8.22 DEFMData Class Reference

Data class for **DEFM** arrays.

```
#include <defm-types.hpp>
```

#### **Public Member Functions**

• DEFMData ()

Vector indicating which covariates are included in the model.

DEFMData (DEFMArray \*array\_, const double \*covariates\_, size\_t obs\_start\_, size\_t X\_ncol\_, size\_t X\_
 nrow )

Constructor.

• double operator() (size t i, size t j) const

Access to the row (i) colum (j) data.

- double at (size\_t i, size\_t j) const
- size\_t ncol () const
- size\_t nrow () const
- void print () const
- ∼DEFMData ()

#### **Public Attributes**

- DEFMArray \* array
- const double \* covariates

Vector of covariates (complete vector)

size\_t obs\_start

Index of the observation in the data.

size\_t X\_ncol

Number of columns in the array of covariates.

size\_t X\_nrow

Number of rows in the array of covariates.

- std::vector< size\_t > covar\_sort
- std::vector< size\_t > covar\_used

Value where the sorting of the covariates is stored.

#### 8.22.1 Detailed Description

Data class for **DEFM** arrays.

This holds information pointing to the data array, including information regarding the number of observations, the time slices of the observation, and the number of covariates in the data.

Definition at line 16 of file defm-types.hpp.

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

• include/barry/models/defm/defm-types.hpp

# 8.23 DEFMRuleData Class Reference

#include <defm-types.hpp>

Inheritance diagram for DEFMRuleData:



#### **Public Member Functions**

- double num (size\_t i) const
- size\_t idx (size\_t i) const
- bool is\_true (size\_t i) const
- DEFMRuleData ()
- DEFMRuleData (std::vector< double > numbers\_, std::vector< size\_t > indices\_, std::vector< bool > logical\_)
- DEFMRuleData (std::vector< double > numbers\_, std::vector< size\_t > indices\_)

## **Public Attributes**

- std::vector< double > numbers
- std::vector< size\_t > indices
- std::vector< bool > logical
- bool init = false

# 8.23.1 Detailed Description

Definition at line 89 of file defm-types.hpp.

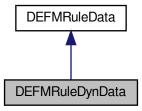
The documentation for this class was generated from the following file:

• include/barry/models/defm/defm-types.hpp

# 8.24 DEFMRuleDynData Class Reference

#include <defm-types.hpp>

Inheritance diagram for DEFMRuleDynData:



Collaboration diagram for DEFMRuleDynData:



# **Public Member Functions**

- DEFMRuleDynData (const std::vector< double > \*counts\_, std::vector< double > numbers\_={}, std::vector< size\_t > indices\_={}, std::vector< bool > logical\_={})
- ∼DEFMRuleDynData ()

## **Public Attributes**

const std::vector< double > \* counts

# 8.24.1 Detailed Description

Definition at line 152 of file defm-types.hpp.

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

include/barry/models/defm/defm-types.hpp

# 8.25 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 (size\_t n)
- ∼Entries ()
- void resize (size\_t n)

# **Public Attributes**

```
• std::vector< size_t > source
```

- std::vector< size\_t > target
- std::vector< Cell\_Type > val

#### 8.25.1 Detailed Description

```
\label{lem:continuous} \begin{split} \text{template} &< \text{typename Cell\_Type} > \\ \text{class Entries} &< \text{Cell\_Type} > \end{split}
```

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

**Template Parameters** 

```
Cell_Type Any type
```

Definition at line 78 of file typedefs.hpp.

#### 8.25.2 Constructor & Destructor Documentation

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

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

Definition at line 84 of file typedefs.hpp.

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

Definition at line 85 of file typedefs.hpp.

# 8.25.2.3 $\sim$ Entries()

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

Definition at line 92 of file typedefs.hpp.

#### 8.25.3 Member Function Documentation

#### 8.25.3.1 resize()

Definition at line 94 of file typedefs.hpp.

# 8.25.4 Member Data Documentation

#### 8.25.4.1 source

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

Definition at line 80 of file typedefs.hpp.

# 8.25.4.2 target

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

Definition at line 81 of file typedefs.hpp.

#### 8.25.4.3 val

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

Definition at line 82 of file typedefs.hpp.

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

• include/barry/typedefs.hpp

#### 8.26 Flock Class Reference

A Flock is a group of Geese.

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

#### **Public Member Functions**

- Flock ()
- ∼Flock ()
- size\_t add\_data (std::vector< std::vector< size\_t > &annotations, std::vector< size\_t > &geneid, std
   ::vector< int > &parent, std::vector< bool > &duplication)

Add a tree to the flock.

• void set seed (const size t &s)

Set the seed of the model.

- void init (size\_t bar\_width=BARRY\_PROGRESS\_BAR\_WIDTH)
- PhyloCounters \* get\_counters ()
- PhyloSupport \* get support fun ()
- std::vector< std::vector< double > > \* get stats support ()
- std::vector< std::vector< double > > \* get\_stats\_target ()
- PhyloModel \* get\_model ()

Returns the joint likelihood of the model.

• Geese \* operator() (size\_t i, bool check\_bounds=true)

Access the i-th geese element.

#### Information about the model

- size\_t nfuns () const noexcept
- size\_t ntrees () const noexcept
- std::vector< size\_t > nnodes () const noexcept
- std::vector< size\_t > nleafs () const noexcept
- size\_t nterms () const
- size\_t support\_size () const noexcept
- std::vector< std::string > colnames () const
- $\bullet \ \ \text{size\_t parse\_polytomies (bool verb=true, std::vector} < \ \text{size\_t} > * \ \text{dist=nullptr) const noexcept} \\$

Check polytomies and return the largest.

void print () const

8.26 Flock Class Reference 149

# **Public Attributes**

- std::vector< Geese > dat
- size\_t nfunctions = 0u
- bool initialized = false
- std::mt19937 rengine
- PhyloModel model = PhyloModel()

# 8.26.1 Detailed Description

A Flock is a group of Geese.

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

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

#### 8.26.2 Constructor & Destructor Documentation

# 8.26.2.1 Flock()

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

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

#### 8.26.2.2 ∼Flock()

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

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

## 8.26.3 Member Function Documentation

## 8.26.3.1 add\_data()

Add a tree to the flock.

#### **Parameters**

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

#### Returns

size\_t The number of tree in the model (starting from zero).

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

#### 8.26.3.2 colnames()

```
std::vector< std::string > Flock::colnames ( ) const [inline]
```

Definition at line 224 of file flock-meat.hpp.

# 8.26.3.3 get\_counters()

```
PhyloCounters * Flock::get_counters ( ) [inline]
```

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

# 8.26.3.4 get\_model()

```
PhyloModel * Flock::get_model ( ) [inline]
```

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

# 8.26.3.5 get\_stats\_support()

```
std::vector< std::vector< double > > * Flock::get_stats_support ( ) [inline]
```

Definition at line 117 of file flock-meat.hpp.

8.26 Flock Class Reference 151

## 8.26.3.6 get\_stats\_target()

```
\verb|std::vector| < \verb|std::vector| < \verb|double| > > * Flock::get_stats_target () | [inline]|
```

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

#### 8.26.3.7 get\_support\_fun()

```
PhyloSupport * Flock::get_support_fun ( ) [inline]
```

Definition at line 110 of file flock-meat.hpp.

# 8.26.3.8 init()

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

#### 8.26.3.9 likelihood\_joint()

Returns the joint likelihood of the model.

#### **Parameters**

par	Vector of model parameters.	
as_log	When true it will return the value as log.	
use_reduced_sequence		
	which is faster.	

## Returns

double

Definition at line 138 of file flock-meat.hpp.

#### 8.26.3.10 nfuns()

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

Definition at line 167 of file flock-meat.hpp.

#### 8.26.3.11 nleafs()

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

Definition at line 195 of file flock-meat.hpp.

## 8.26.3.12 nnodes()

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

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

# 8.26.3.13 nterms()

```
size_t Flock::nterms ( ) const [inline]
```

Definition at line 209 of file flock-meat.hpp.

## 8.26.3.14 ntrees()

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

Definition at line 174 of file flock-meat.hpp.

# 8.26.3.15 operator()()

Access the i-th geese element.

8.26 Flock Class Reference 153

#### **Parameters**

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

Returns

Geese \*

Definition at line 302 of file flock-meat.hpp.

# 8.26.3.16 parse\_polytomies()

Check polytomies and return the largest.

Definition at line 231 of file flock-meat.hpp.

## 8.26.3.17 print()

```
void Flock::print ( ) const [inline]
```

Definition at line 258 of file flock-meat.hpp.

# 8.26.3.18 set\_seed()

Set the seed of the model.

**Parameters** 

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

Definition at line 42 of file flock-meat.hpp.

# 8.26.3.19 support\_size()

```
size_t Flock::support_size ( ) const [inline], [noexcept]
```

Definition at line 217 of file flock-meat.hpp.

# 8.26.4 Member Data Documentation

# 8.26.4.1 dat

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

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

# 8.26.4.2 initialized

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

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

#### 8.26.4.3 model

```
PhyloModel Flock::model = PhyloModel()
```

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

# 8.26.4.4 nfunctions

```
size_t Flock::nfunctions = 0u
```

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

#### 8.26.4.5 rengine

```
std::mt19937 Flock::rengine
```

Definition at line 22 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-meat.hpp

# 8.27 FreqTable < T > Class Template Reference

Frequency table of vectors.

```
#include <freqtable.hpp>
```

#### **Public Member Functions**

```
• FreqTable ()
```

- ∼FreqTable ()
- size\_t add (const std::vector< T > &x, size\_t \*h\_precomp)
- Counts\_type as\_vector () const
- const std::vector< double > & get\_data () const
- const std::unordered\_map< size\_t, size\_t > & get\_index () const
- void clear ()
- void reserve (size\_t n, size\_t k)
- void print () const
- · size\_t size () const noexcept

Number of unique elements in the table. (.

size\_t make\_hash (const std::vector< T > &x) const

# 8.27.1 Detailed Description

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

Frequency table of vectors.

This is mostly used in Support. The main data is contained in the data double vector. The matrix is stored in a row-wise fashion, where the first element is the frequency with which the vector is observed.

For example, in a model with k terms the first k + 1 elements of data would be:

- · weights
- term 1
- term 2
- ...
- · term k

Definition at line 22 of file freqtable.hpp.

# 8.27.2 Constructor & Destructor Documentation

# 8.27.2.1 FreqTable()

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

Definition at line 34 of file freqtable.hpp.

#### 8.27.2.2 ∼FreqTable()

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

Definition at line 35 of file freqtable.hpp.

#### 8.27.3 Member Function Documentation

# 8.27.3.1 add()

Definition at line 59 of file freqtable.hpp.

# 8.27.3.2 as\_vector()

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

Definition at line 139 of file freqtable.hpp.

#### 8.27.3.3 clear()

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

Definition at line 168 of file freqtable.hpp.

#### 8.27.3.4 get\_data()

```
template<typename T = double>
const std::vector< double >& FreqTable< T >::get_data ( ) const [inline]
```

Definition at line 40 of file freqtable.hpp.

# 8.27.3.5 get\_index()

```
template<typename T = double>
const std::unordered_map<size_t,size_t>& FreqTable< T >::get_index ( ) const [inline]
```

Definition at line 41 of file freqtable.hpp.

#### 8.27.3.6 make\_hash()

Definition at line 239 of file freqtable.hpp.

# 8.27.3.7 print()

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

Definition at line 204 of file freqtable.hpp.

#### 8.27.3.8 reserve()

Definition at line 182 of file freqtable.hpp.

#### 8.27.3.9 size()

```
template<typename T >
size_t FreqTable< T >::size [inline], [noexcept]
Number of unique elements in the table. (.
```

Returns

size t

Definition at line 231 of file freqtable.hpp.

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

• include/barry/freqtable.hpp

#### 8.28 Geese Class Reference

Annotated Phylo Model.

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

#### **Public Member Functions**

- ~Geese ()
- void init (size t bar width=BARRY PROGRESS BAR WIDTH)
- void inherit support (const Geese &model, bool delete support =false)
- void calc\_sequence (Node \*n=nullptr)
- void calc reduced sequence ()
- double likelihood (const std::vector< double > &par, bool as\_log=false, bool use\_reduced\_sequence=true)
- double likelihood exhaust (const std::vector< double > &par)
- std::vector< double > get\_probabilities () const
- void set\_seed (const size\_t &s)
- std::vector< std::vector< size\_t >> simulate (const std::vector< double > &par)
- std::vector< std::vector< double > > observed\_counts ()
- void print\_observed\_counts ()
- · void print () const

Prints information about the GEESE.

- void init node (Node &n)
- void update\_annotations (size\_t nodeid, std::vector< size\_t > newann)
- std::vector< std::vector< bool >> get\_states () const

Powerset of a gene's possible states.

std::vector < size\_t > get\_annotated\_nodes () const

Returns the ids of the nodes with at least one annotation.

#### Construct a new Geese object

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

#### **Parameters**

annotations	A vector of vectors with annotations. It should be of length $k$ (number of functions). Each vector should be of length $N$ (equal to the number of nodes, including interior). Possible values are 0, 1, and 9.	
geneid	Id of the gene. It should be of length N.	
parent	Id of the parent gene. Also of length ${\tt N}$	
duplication	Logical scalar indicating the type of event (true: duplication, false: speciation.)	

The ordering of the entries does not matter. Passing the nodes in post order or not makes no difference to the constructor.

- Geese ()
- Geese (std::vector< std::vector< size\_t > & annotations, std::vector< size\_t > & 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

#### **Parameters**

verb When true it will print out information about the er	ncountered polytomies.
---	------------------------

• size\_t nfuns () const noexcept

Number of functions analyzed.

• size\_t nnodes () const noexcept

Number of nodes (interior + leaf)

• size\_t nleafs () const noexcept

Number of leaf.

• size\_t nterms () const

Number of terms included.

size\_t support\_size () const noexcept

Number of unique sets of sufficient stats.

std::vector< size\_t > nannotations () const noexcept

Number of annotations.

• std::vector< std::string > colnames () const

Names of the terms in the model.

size\_t parse\_polytomies (bool verb=true, std::vector< size\_t > \*dist=nullptr) const noexcept
 Check polytomies and return the largest.

#### **Geese prediction**

Calculate the conditional probability

## **Parameters**

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

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

#### Returns

std::vector< double > Returns the posterior probability

- std::vector< std::vector< double >> predict (const std::vector< double > &par, std::vector< std::vector< double >> \*res\_prob=nullptr, bool leave\_one\_out=false, bool only\_annotated=false, bool use\_reduced = sequence=true)
- std::vector < std::vector < double > > predict\_backend (const std::vector < double > &par, bool use\_←
  reduced\_sequence, const std::vector < size\_t > &preorder)
- std::vector< std::vector< double > > predict\_exhaust\_backend (const std::vector< double > &par, const std::vector< size t > &preorder)
- std::vector< std::vector< double > > predict\_exhaust (const std::vector< double > &par)
- std::vector< std::vector< double > > predict\_sim (const std::vector< double > &par, bool only\_

   annotated=false, size\_t nsims=10000u)

#### Non-const pointers to shared objects in <tt>Geese</tt>

These functions provide direct access to some member objects that are shared by the nodes within Geese.

#### Returns

```
get_rengine() returns the Pseudo-RNG engine used.
get_counters() returns the vector of counters used.
get_model() returns the Model object used.
get_support_fun() returns the computed support of the model.
```

- std::mt19937 \* get\_rengine ()
- PhyloCounters \* get\_counters ()
- PhyloModel \* get\_model ()
- PhyloSupport \* get\_support\_fun ()

#### **Public Attributes**

- · size\_t nfunctions
- std::map< size\_t, Node > nodes
- barry::MapVec\_type< size\_t > map\_to\_nodes
- $std::vector < std::vector < size_t >>> pset_loc$

Locations of columns.

- std::vector< size t > sequence
- std::vector< size\_t > reduced\_sequence
- bool initialized = false
- bool delete\_rengine = false
- bool delete\_support = false

#### **Static Public Attributes**

```
static const size_t etype_default = 1ul
```

- static const size\_t etype\_speciation = 0ul
- static const size\_t etype\_duplication = 1ul
- static const size\_t etype\_either = 2ul

8.28 Geese Class Reference 161

## 8.28.1 Detailed Description

Annotated Phylo Model.

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

Class representing a phylogenetic tree model with annotations.

The Geese class represents a phylogenetic tree model with annotations. It includes a total of N+1 nodes, the +1 being the root node. The class provides methods for initializing the model, calculating the likelihood, simulating trees, and making predictions.

The class includes shared objects within a Geese object, such as rengine, model, states, n\_zeros, n\_ $\leftarrow$  ones, n\_dupl\_events, and n\_spec\_events. It also includes information about the type of event, such as etype\_default, etype\_speciation, etype\_duplication, and etype\_either.

The class provides constructors, a destructor, and methods for initializing the model, inheriting support, calculating the sequence, calculating the reduced sequence, calculating the likelihood, calculating the likelihood exhaustively, getting probabilities, setting the seed, simulating trees, parsing polytomies, getting observed counts, printing information about the GEESE, and making predictions.

See also

Flock

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

## 8.28.2 Constructor & Destructor Documentation

#### 8.28.2.1 Geese() [1/4]

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

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

#### 8.28.2.2 Geese() [2/4]

```
Geese::Geese (
          std::vector< std::vector< size_t > & annotations,
          std::vector< size_t > & geneid,
          std::vector< int > & parent,
          std::vector< bool > & duplication ) [inline]
```

Definition at line 20 of file geese-meat-constructors.hpp.

#### 8.28.2.3 Geese() [3/4]

Definition at line 216 of file geese-meat-constructors.hpp.

# 8.28.2.4 Geese() [4/4]

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

#### 8.28.2.5 ∼Geese()

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

Definition at line 144 of file geese-meat.hpp.

# 8.28.3 Member Function Documentation

## 8.28.3.1 calc\_reduced\_sequence()

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

Definition at line 383 of file geese-meat.hpp.

# 8.28.3.2 calc\_sequence()

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

Definition at line 339 of file geese-meat.hpp.

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#### 8.28.3.3 colnames()

```
std::vector< std::string > Geese::colnames ( ) const [inline]
```

Names of the terms in the model.

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

#### 8.28.3.4 get\_annotated\_nodes()

```
std::vector< size_t > Geese::get_annotated_nodes ( ) const [inline]
```

Returns the ids of the nodes with at least one annotation.

Definition at line 721 of file geese-meat.hpp.

#### 8.28.3.5 get\_counters()

```
PhyloCounters * Geese::get_counters ( ) [inline]
```

Definition at line 704 of file geese-meat.hpp.

# 8.28.3.6 get\_model()

```
PhyloModel * Geese::get_model ( ) [inline]
```

Definition at line 709 of file geese-meat.hpp.

## 8.28.3.7 get\_probabilities()

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

Definition at line 431 of file geese-meat.hpp.

# 8.28.3.8 get\_rengine()

```
std::mt19937 * Geese::get_rengine ( ) [inline]
```

Definition at line 699 of file geese-meat.hpp.

#### 8.28.3.9 get\_states()

```
std::vector< std::vector< bool > > Geese::get_states ( ) const [inline]
```

Powerset of a gene's possible states.

This list of vectors is used throughout Geese. It lists all possible combinations of functional states for any gene. Thus, for P functions, there will be  $2^{P}$  possible combinations.

Returns

```
std::vector< std::vector< bool >> of length 2^{^{\text{P}}}.
```

Definition at line 717 of file geese-meat.hpp.

#### 8.28.3.10 get\_support\_fun()

```
PhyloSupport * Geese::get_support_fun ( ) [inline]
```

Definition at line 713 of file geese-meat.hpp.

#### 8.28.3.11 inherit\_support()

Definition at line 282 of file geese-meat.hpp.

## 8.28.3.12 init()

Definition at line 156 of file geese-meat.hpp.

#### 8.28.3.13 init\_node()

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

#### 8.28.3.14 likelihood()

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

#### 8.28.3.15 likelihood\_exhaust()

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

#### 8.28.3.16 nannotations()

```
std::vector< size_t > Geese::nannotations ( ) const [inline], [noexcept]
```

Number of annotations.

Definition at line 496 of file geese-meat.hpp.

# 8.28.3.17 nfuns()

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

Number of functions analyzed.

Definition at line 452 of file geese-meat.hpp.

#### 8.28.3.18 nleafs()

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

Number of leaf.

Definition at line 466 of file geese-meat.hpp.

#### 8.28.3.19 nnodes()

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

Number of nodes (interior + leaf)

Definition at line 459 of file geese-meat.hpp.

#### 8.28.3.20 nterms()

```
size_t Geese::nterms ( ) const [inline]
```

Number of terms included.

Definition at line 478 of file geese-meat.hpp.

#### 8.28.3.21 observed\_counts()

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

Definition at line 547 of file geese-meat.hpp.

# 8.28.3.22 operator=() [1/2]

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

# 8.28.3.24 parse\_polytomies()

Check polytomies and return the largest.

Definition at line 512 of file geese-meat.hpp.

#### 8.28.3.25 predict()

Definition at line 272 of file geese-meat-predict.hpp.

# 8.28.3.26 predict\_backend()

< True if the array belongs to the set

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

#### 8.28.3.27 predict\_exhaust()

Definition at line 5 of file geese-meat-predict exhaust.hpp.

#### 8.28.3.28 predict\_exhaust\_backend()

Definition at line 47 of file geese-meat-predict exhaust.hpp.

#### 8.28.3.29 predict\_sim()

```
std::vector< std::vector< double > > Geese::predict_sim (
    const std::vector< double > & par,
    bool only_annotated = false,
    size_t nsims = 10000u ) [inline]
```

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

## 8.28.3.30 print()

```
void Geese::print ( ) const [inline]
```

Prints information about the GEESE.

Definition at line 681 of file geese-meat.hpp.

# 8.28.3.31 print\_observed\_counts()

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

Definition at line 618 of file geese-meat.hpp.

#### 8.28.3.32 set\_seed()

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

## 8.28.3.33 simulate()

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

## 8.28.3.34 support\_size()

```
size_t Geese::support_size ( ) const [inline], [noexcept]
```

Number of unique sets of sufficient stats.

Definition at line 486 of file geese-meat.hpp.

#### 8.28.3.35 update\_annotations()

Definition at line 310 of file geese-meat.hpp.

#### 8.28.4 Member Data Documentation

## 8.28.4.1 delete\_rengine

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

Definition at line 142 of file geese-bones.hpp.

# 8.28.4.2 delete\_support

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

Definition at line 143 of file geese-bones.hpp.

# 8.28.4.3 etype\_default

```
const size_t Geese::etype_default = 1ul [static]
```

Definition at line 156 of file geese-bones.hpp.

#### 8.28.4.4 etype\_duplication

```
const size_t Geese::etype_duplication = 1ul [static]
```

Definition at line 158 of file geese-bones.hpp.

# 8.28.4.5 etype\_either

```
const size_t Geese::etype_either = 2ul [static]
```

Definition at line 159 of file geese-bones.hpp.

#### 8.28.4.6 etype\_speciation

```
const size_t Geese::etype_speciation = Oul [static]
```

Definition at line 157 of file geese-bones.hpp.

#### 8.28.4.7 initialized

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

Definition at line 141 of file geese-bones.hpp.

# 8.28.4.8 map\_to\_nodes

```
barry::MapVec_type< size_t > Geese::map_to_nodes
```

Definition at line 133 of file geese-bones.hpp.

#### 8.28.4.9 nfunctions

```
size_t Geese::nfunctions
```

Definition at line 131 of file geese-bones.hpp.

#### 8.28.4.10 nodes

```
std::map< size_t, Node > Geese::nodes
```

Definition at line 132 of file geese-bones.hpp.

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#### 8.28.4.11 pset\_loc

```
std::vector< std::vector< size_t > > > Geese::pset_loc
```

Locations of columns.

Definition at line 134 of file geese-bones.hpp.

#### 8.28.4.12 reduced\_sequence

```
std::vector< size_t > Geese::reduced_sequence
```

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

#### 8.28.4.13 sequence

```
std::vector< size_t > Geese::sequence
```

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

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

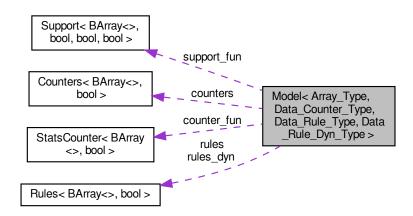
- include/barry/models/geese/geese-bones.hpp
- include/barry/models/geese/geese-meat-constructors.hpp
- include/barry/models/geese/geese-meat-likelihood.hpp
- include/barry/models/geese/geese-meat-likelihood exhaust.hpp
- include/barry/models/geese/geese-meat-predict.hpp
- include/barry/models/geese/geese-meat-predict\_exhaust.hpp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

# 8.29 Model < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_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, Data\_Rule\_Dyn\_Type >:



#### **Public Member Functions**

- void set\_rengine (std::mt19937 \*rengine\_, bool delete\_=false)
- void set\_seed (size\_t s)
- Model ()
- Model (size\_t size\_)
- Model (const Model< Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type > &Model )
- Model < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type > & operator= (const Model < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type > & Model\_)
- virtual ∼Model ()
- void store\_psets () noexcept
- std::vector< double > gen\_key (const Array\_Type &Array\_)
- size\_t add\_array (const Array\_Type &Array\_, bool force\_new=false)

Adds an array to the support of not already included.

- void print\_stats (size\_t i) const
- · virtual void print () const

Prints information about the model.

- Array\_Type sample (const Array\_Type &Array\_, const std::vector< double > &params={})
- Array\_Type sample (const size\_t &i, const std::vector< double > &params)
- double conditional\_prob (const Array\_Type &Array\_, const std::vector< double > &params, size\_t i, size\_t j)
   Conditional probability ("Gibbs sampler")
- const std::mt19937 \* get rengine () const
- Counters < Array\_Type, Data\_Counter\_Type > \* get\_counters ()

- Rules< Array\_Type, Data\_Rule\_Type > \* get\_rules ()
- Rules< Array\_Type, Data\_Rule\_Dyn\_Type > \* get\_rules\_dyn ()
- Support < Array Type, Data Counter Type, Data Rule Type, Data Rule Dyn Type > \* get support fun ()

#### 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\_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)
- void set\_counters (Counters < Array\_Type, Data\_Counter\_Type > \*counters\_)
- void add hasher (Hasher fun type< Array Type, Data Counter Type > fun )

#### 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\_fun\_type< Array\_Type, Data\_Rule\_Type > count\_fun\_, Data\_Rule\_Type data\_)
- void set\_rules (Rules < Array\_Type, Data\_Rule\_Type > \*rules\_)
- void add rule dyn (Rule < Array Type, Data Rule Dyn Type > &rule)
- void set\_rules\_dyn (Rules< Array\_Type, Data\_Rule\_Dyn\_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 size t &i, bool as log=false)
- double likelihood (const std::vector< double > &params, const std::vector< double > &target\_, const size t &i, bool as log=false)
- double likelihood (const std::vector< double > &params, const double \*target\_, const size\_t &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 size t &i, bool as log=false)
- const std::vector< Array\_Type > \* get\_pset (const size\_t &i)
- const std::vector< double > \* get\_pset\_stats (const size\_t &i)

#### Size of the model

Number of different supports included in the model

This will return the size of stats\_target.

#### 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.
```

- size t size () const noexcept
- size\_t size\_unique () const noexcept
- size\_t nterms () const noexcept
- size\_t nrules () const noexcept
- size\_t nrules\_dyn () const noexcept
- size t support size () const noexcept
- std::vector< std::string > colnames () const
- std::vector< std::vector< double >>\* get\_stats\_target ()

Raw pointers to the support and target statistics.

- std::vector< std::vector< double >> \* get\_stats\_support ()
- std::vector< size\_t > \* get\_arrays2support ()
- std::vector< std::vector< Array\_Type > > \* get\_pset\_arrays ()
- $std::vector < std::vector < double >> * get_pset_stats ()$

Statistics of the support(s)

- std::vector< std::vector< double > > \* get\_pset\_probs ()
- void set\_transform\_model (std::function< std::vector< double >(double \*, size\_t)> fun, std::vector< std

   ::string > names)

Set the transform\_model\_fun object.

std::vector< double > transform\_model (double \*data, size\_t k)

#### **Protected Attributes**

• MapVec\_type< double, size\_t > 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
- bool delete\_counters = false
- bool delete rules = false
- bool delete\_rules\_dyn = false
- std::function< std::vector< double >double \*, size\_t k)> transform\_model\_fun = nullptr

Transformation of the model.

• std::vector< std::string > transform\_model\_term\_names

#### Random number generation

Random number generation

- std::mt19937 \* rengine = nullptr
- bool delete rengine = false

### Information about the arrays used in the model

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

Each vector of stats\_support has the data stored in a row-wise order, with each row starting with the weights, e.g., in a model with k terms the first k + 1 elements of stats\_support would be:

- · weights
- term 1
- term 2
- ...
- term k
- std::vector < std::vector < double > > stats\_support
   Sufficient statistics of the model (support)
- std::vector< size t > stats support n arrays

Number of arrays included per support.

std::vector< std::vector< double >> stats\_target

Target statistics of the model.

std::vector< size\_t > arrays2support

### Container space for the powerset (and its sufficient stats\_target)

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

Arrays of the support(s)

std::vector< std::vector< double >> pset\_stats

Statistics of the support(s)

std::vector< std::vector< double >> pset probs

Probabilities of the support(s)

## Functions to compute statistics

Arguments are recycled to save memory and computation.

- $\bullet \ \ Counters < Array\_Type, \ Data\_Counter\_Type > * counters$
- Rules < Array\_Type, Data\_Rule\_Type > \* rules
- Rules < Array\_Type, Data\_Rule\_Dyn\_Type > \* rules\_dyn
- Support < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type > support\_fun
- StatsCounter< Array\_Type, Data\_Counter\_Type > counter\_fun

### 8.29.1 Detailed Description

template < typename Array\_Type = BArray <>>, typename Data\_Counter\_Type = bool, typename Data\_Rule\_Type = bool, typename Data\_Rule\_Dyn\_Type = bool>

class Model < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type >

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

$$\frac{\exp\left(\theta^{\mathsf{t}}c(A)\right)}{\sum_{A'\in\mathcal{A}}\exp\left(\theta^{\mathsf{t}}c(A')\right)}$$

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 34 of file model-bones.hpp.

### 8.29.2 Constructor & Destructor Documentation

## 8.29.2.1 Model() [1/3]

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

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

#### 8.29.2.2 Model() [2/3]

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

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

Definition at line 204 of file model-meat.hpp.

### 8.29.2.4 ∼Model()

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

Definition at line 159 of file model-bones.hpp.

### 8.29.3 Member Function Documentation

### 8.29.3.1 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.

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

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

### 8.29.3.4 add\_hasher()

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

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::add_rule (
Rule< Array_Type, Data_Rule_Type > & rule )
```

#### 8.29.3.6 add rule() [2/2]

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::add_rule (
    Rule_fun_type< Array_Type, Data_Rule_Type > count_fun_,
    Data_Rule_Type data_ )
```

### 8.29.3.7 add\_rule\_dyn() [1/2]

#### 8.29.3.8 add\_rule\_dyn() [2/2]

#### 8.29.3.9 colnames()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::string > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_\times
Dyn_Type >::colnames () const
```

### 8.29.3.10 conditional\_prob()

Conditional probability ("Gibbs sampler")

Computes the conditional probability of observing  $P\{Y(i,j) = | Y^C, \text{ theta}\}$ , i.e., the probability of observing the entry Y(i,j) equal to one given the rest of the array.

#### **Parameters**

Array←	Array to check
_	
params	Vector of parameters
i	Row entry
j	Column entry

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#### Returns

double The conditional probability

### 8.29.3.11 gen\_key()

### 8.29.3.12 get\_arrays2support()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< size_t >* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_\times
Type >::get_arrays2support ()
```

### 8.29.3.13 get\_counters()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Counters<Array_Type, Data_Counter_Type>* Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_counters ()
```

#### 8.29.3.14 get norm const()

#### 8.29.3.15 get pset()

#### 8.29.3.16 get\_pset\_arrays()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector< Array_Type > >* Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_pset_arrays ()
```

#### 8.29.3.17 get\_pset\_probs()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector<double> >* Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_pset_probs ()
```

## 8.29.3.18 get\_pset\_stats() [1/2]

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector<double> >* Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_pset_stats ()
```

Statistics of the support(s)

### 8.29.3.19 get\_pset\_stats() [2/2]

## 8.29.3.20 get\_rengine()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
const std::mt19937* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type
>::get_rengine ( ) const
```

#### 8.29.3.21 get\_rules()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Rules<Array_Type, Data_Rule_Type>* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data←
_Rule_Dyn_Type >::get_rules ()
```

#### 8.29.3.22 get\_rules\_dyn()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Rules<Array_Type, Data_Rule_Dyn_Type>* Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_rules_dyn ()
```

### 8.29.3.23 get\_stats\_support()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector< double > >* Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_stats_support ()
```

#### 8.29.3.24 get\_stats\_target()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarray_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector< double > >* Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_stats_target ()
```

Raw pointers to the support and target statistics.

The support of the model is stored as a vector of vector < double>. Each element of it contains the support for an specific type of array included. It represents an array of size  $(k + 1) \times n \text{ unique elements}$ , with the data stored by-row. The last element of each entry corresponds to the weights, i.e., the frequency with which such sufficient statistics are observed in the support.

### 8.29.3.25 get\_support\_fun()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Support<Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type>* Model< Array_Type,
Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::get_support_fun ()
```

### 8.29.3.26 likelihood() [1/4]

### 8.29.3.27 likelihood() [2/4]

### 8.29.3.28 likelihood() [3/4]

#### 8.29.3.29 likelihood() [4/4]

#### 8.29.3.30 likelihood\_total()

#### 8.29.3.31 nrules()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::nrules ()
const [noexcept]
```

## 8.29.3.32 nrules\_dyn()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::nrules_dyn
( ) const [noexcept]
```

#### 8.29.3.33 nterms()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::nterms ()
const [noexcept]
```

#### 8.29.3.34 operator=()

Definition at line 248 of file model-meat.hpp.

#### 8.29.3.35 print()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::print [inline],
[virtual]
```

Prints information about the model.

Definition at line 971 of file model-meat.hpp.

#### 8.29.3.36 print\_stats()

### 8.29.3.37 sample() [1/2]

## 8.29.3.38 sample() [2/2]

Definition at line 1098 of file model-meat.hpp.

#### 8.29.3.39 set counters()

#### 8.29.3.40 set\_rengine()

Definition at line 129 of file model-bones.hpp.

## 8.29.3.41 set\_rules()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::set_rules (
Rules< Array_Type, Data_Rule_Type > * rules_ )
```

#### 8.29.3.42 set rules dyn()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::set_rules_dyn
(
Rules< Array_Type, Data_Rule_Dyn_Type > * rules_ )
```

### 8.29.3.43 set\_seed()

Definition at line 139 of file model-bones.hpp.

### 8.29.3.44 set\_transform\_model()

Set the transform model fun object.

The transform\_model function is used to transform the data

#### **Parameters**

data	
target	
n_arrays	
arrays2support	

## 8.29.3.45 size()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::size ()
const [noexcept]
```

#### 8.29.3.46 size\_unique()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::size_unique
( ) const [noexcept]
```

## 8.29.3.47 store\_psets()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::store_psets (
) [noexcept]
```

#### 8.29.3.48 support\_size()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::support_←
size ( ) const [noexcept]
```

#### 8.29.3.49 transform\_model()

### 8.29.4 Member Data Documentation

## 8.29.4.1 arrays2support

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< size_t > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::arrays2support [protected]
```

Definition at line 65 of file model-bones.hpp.

#### 8.29.4.2 counter\_fun

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
StatsCounter<Array_Type, Data_Counter_Type> Model< Array_Type, Data_Counter_Type, Data_Rule_Dyn_Type >::counter_fun [protected]
```

Definition at line 95 of file model-bones.hpp.

#### 8.29.4.3 counters

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Counters<Array_Type, Data_Counter_Type>* Model< Array_Type, Data_Counter_Type, Data_Rule_Dyn_Type >::counters [protected]
```

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

#### 8.29.4.4 delete\_counters

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_←
counters = false [protected]
```

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

#### 8.29.4.5 delete rengine

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_←
rengine = false [protected]
```

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

#### 8.29.4.6 delete\_rules

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_rules
= false [protected]
```

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

### 8.29.4.7 delete\_rules\_dyn

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_←
rules_dyn = false [protected]
```

Definition at line 105 of file model-bones.hpp.

### 8.29.4.8 first\_calc\_done

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

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

### 8.29.4.9 keys2support

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
MapVec_type< double, size_t > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_←
Rule_Dyn_Type >::keys2support [protected]
```

Map of types of arrays to support sets.

This is of the same length as the vector stats\_target.

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

### 8.29.4.10 normalizing constants

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< double > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::normalizing_constants [protected]
```

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

## 8.29.4.11 params\_last

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

Vector of the previously used parameters.

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

## 8.29.4.12 pset\_arrays

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector< Array_Type >> Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::pset_arrays [protected]
```

Arrays of the support(s)

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

#### 8.29.4.13 pset probs

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

Probabilities of the support(s)

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

## 8.29.4.14 pset\_stats

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

Statistics of the support(s)

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

#### 8.29.4.15 rengine

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

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

#### 8.29.4.16 rules

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Rules<Array_Type, Data_Rule_Type>* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data←
_Rule_Dyn_Type >::rules [protected]
```

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

### 8.29.4.17 rules\_dyn

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Rules<Array_Type, Data_Rule_Dyn_Type>* Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::rules_dyn [protected]
```

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

#### 8.29.4.18 stats support

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector< double > > Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::stats_support [protected]
```

Sufficient statistics of the model (support)

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

#### 8.29.4.19 stats\_support\_n\_arrays

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< size_t > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::stats_support_n_arrays [protected]
```

Number of arrays included per support.

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

## 8.29.4.20 stats\_target

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector< double > > Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::stats_target [protected]
```

Target statistics of the model.

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

#### 8.29.4.21 support fun

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

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

### 8.29.4.22 transform\_model\_fun

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::function<std::vector<double>double *, size_t k)> Model< Array_Type, Data_Counter_Type,
Data_Rule_Type, Data_Rule_Dyn_Type >::transform_model_fun = nullptr [protected]
```

Transformation of the model.

When specified, this function will update the model by modifying the linear equation. For example, if the user wanted to add interaction terms, rescale, or apply other operations of the sorts, the user can do such through this function.

The function should return void and receive the following arguments:

- data Pointer to the first element of the set of sufficient statistics
- k size\_t indicating the number of sufficient statistics

#### Returns

Nothing, but it will modify the model data.

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

### 8.29.4.23 transform\_model\_term\_names

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::string > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_\times
Dyn_Type >::transform_model_term_names [protected]
```

Definition at line 125 of file model-bones.hpp.

#### 8.29.4.24 with\_pset

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::with_pset =
false [protected]
```

Definition at line 80 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

### 8.30 NetCounterData Class Reference

Data class used to store arbitrary size\_t or double vectors.

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

### **Public Member Functions**

- NetCounterData ()
- NetCounterData (const std::vector < size\_t > indices\_, const std::vector < double > numbers\_)
- ∼NetCounterData ()

#### **Public Attributes**

- std::vector< size t > indices
- std::vector< double > numbers

## 8.30.1 Detailed Description

Data class used to store arbitrary size\_t or double vectors.

Definition at line 56 of file network.hpp.

#### 8.30.2 Constructor & Destructor Documentation

### 8.30.2.1 NetCounterData() [1/2]

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

Definition at line 62 of file network.hpp.

### 8.30.2.2 NetCounterData() [2/2]

Definition at line 63 of file network.hpp.

### 8.30.2.3 ~NetCounterData()

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

Definition at line 68 of file network.hpp.

#### 8.30.3 Member Data Documentation

### 8.30.3.1 indices

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

Definition at line 59 of file network.hpp.

### 8.30.3.2 numbers

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

Definition at line 60 of file network.hpp.

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

• include/barry/counters/network.hpp

### 8.31 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.

 $\bullet \ \ {\tt NetworkData} \ ({\tt std::vector} < {\tt std::vector} < {\tt double} > > {\tt vertex\_attr\_}, \ {\tt bool} \ {\tt directed\_=true}) \\$ 

Constructor using multiple attributes.

∼NetworkData ()

### **Public Attributes**

- bool directed = true
- std::vector< std::vector< double > > vertex attr

## 8.31.1 Detailed Description

Data class for Networks.

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

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

Definition at line 19 of file network.hpp.

### 8.31.2 Constructor & Destructor Documentation

### 8.31.2.1 NetworkData() [1/3]

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

Definition at line 25 of file network.hpp.

### 8.31.2.2 NetworkData() [2/3]

Constructor using a single attribute.

#### **Parameters**

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

Definition at line 33 of file network.hpp.

### 8.31.2.3 NetworkData() [3/3]

Constructor using multiple attributes.

#### **Parameters**

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

Definition at line 45 of file network.hpp.

### 8.31.2.4 ~NetworkData()

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

Definition at line 51 of file network.hpp.

### 8.31.3 Member Data Documentation

### 8.31.3.1 directed

```
bool NetworkData::directed = true
```

Definition at line 22 of file network.hpp.

### 8.31.3.2 vertex\_attr

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

Definition at line 23 of file network.hpp.

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

• include/barry/counters/network.hpp

# 8.32 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
- size\_t noffspring () const noexcept
- bool is\_leaf () const noexcept

## Construct a new Node object

- Node ()
- Node (size\_t id\_, size\_t ord\_, bool duplication\_)
- Node (size\_t id\_, size\_t ord\_, std::vector < size\_t > annotations\_, bool duplication\_)
- Node (Node &&x) noexcept
- Node (const Node &x)

8.32 Node Class Reference 199

#### **Public Attributes**

```
• size tid
```

Id of the node (as specified in the input)

· size tord

Order in which the node was created.

PhyloArray array

Array of the node.

std::vector< size\_t > annotations

Observed annotations (only defined for Geese)

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

Arrays given all possible states.

• std::vector< bool > arrays\_valid = {}

Whether the arrays are valid according to the rules of the model.

• Node \* parent = nullptr

Parent node.

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

Offspring nodes.

std::vector< size\_t > 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.

### 8.32.1 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.

### 8.32.2 Constructor & Destructor Documentation

#### 8.32.2.1 Node() [1/5]

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

Definition at line 39 of file geese-node-bones.hpp.

### 8.32.2.2 Node() [2/5]

Definition at line 59 of file geese-node-bones.hpp.

#### 8.32.2.3 Node() [3/5]

Definition at line 65 of file geese-node-bones.hpp.

### 8.32.2.4 Node() [4/5]

Definition at line 72 of file geese-node-bones.hpp.

## 8.32.2.5 Node() [5/5]

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

### 8.32.2.6 ∼Node()

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

Definition at line 50 of file geese-node-bones.hpp.

## 8.32.3 Member Function Documentation

8.32 Node Class Reference 201

### 8.32.3.1 get\_parent()

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

Definition at line 104 of file geese-node-bones.hpp.

### 8.32.3.2 is\_leaf()

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

Definition at line 116 of file geese-node-bones.hpp.

### 8.32.3.3 noffspring()

```
size_t Node::noffspring ( ) const [inline], [noexcept]
```

Definition at line 110 of file geese-node-bones.hpp.

#### 8.32.4 Member Data Documentation

### 8.32.4.1 annotations

```
std::vector< size_t > Node::annotations
```

Observed annotations (only defined for Geese)

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

#### 8.32.4.2 array

PhyloArray Node::array

Array of the node.

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

### 8.32.4.3 arrays

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

Arrays given all possible states.

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

### 8.32.4.4 arrays\_valid

```
std::vector< bool > Node::arrays_valid = {}
```

Whether the arrays are valid according to the rules of the model.

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

### 8.32.4.5 duplication

```
bool Node::duplication
```

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

### 8.32.4.6 id

```
size_t Node::id
```

Id of the node (as specified in the input)

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

#### 8.32.4.7 narray

```
std::vector< size_t > Node::narray = {}
```

ID of the array in the model.

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

8.32 Node Class Reference 203

### 8.32.4.8 offspring

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

Offspring nodes.

Definition at line 26 of file geese-node-bones.hpp.

#### 8.32.4.9 ord

```
size_t Node::ord
```

Order in which the node was created.

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

### 8.32.4.10 parent

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

Parent node.

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

### 8.32.4.11 probability

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

The probability of observing each state.

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

## 8.32.4.12 subtree\_prob

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

Induced subtree probabilities.

Definition at line 30 of file geese-node-bones.hpp.

#### 8.32.4.13 visited

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

Definition at line 28 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

### 8.33 NodeData Class Reference

Data definition for the PhyloArray class.

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

#### **Public Member Functions**

NodeData (const std::vector< double > &blengths\_, const std::vector< bool > &states\_, bool duplication
 —=true)

#### **Public Attributes**

```
    std::vector< double > blengths = {}
    std::vector< bool > states = {}
    bool duplication = true
```

## 8.33.1 Detailed Description

Data definition for the PhyloArray class.

This holds basic information about a given node.

Definition at line 12 of file geese-types.hpp.

### 8.33.2 Constructor & Destructor Documentation

#### 8.33.2.1 NodeData()

Definition at line 32 of file geese-types.hpp.

#### 8.33.3 Member Data Documentation

#### 8.33.3.1 blengths

```
std::vector< double > NodeData::blengths = {}
```

Branch length.

Definition at line 18 of file geese-types.hpp.

#### 8.33.3.2 duplication

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

Definition at line 28 of file geese-types.hpp.

#### 8.33.3.3 states

```
std::vector< bool > NodeData::states = {}
```

State of the parent node.

Definition at line 23 of file geese-types.hpp.

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

• include/barry/models/geese/geese-types.hpp

# 8.34 PhyloCounterData Class Reference

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

### **Public Member Functions**

```
    PhyloCounterData (std::vector< size_t > data_, std::vector< double > *counters_=nullptr)
```

- PhyloCounterData ()
- size\_t at (size\_t d)
- size\_t operator() (size\_t d)
- size\_t operator[] (size\_t d)
- void reserve (size\_t x)
- void push\_back (size\_t x)
- void shrink\_to\_fit ()
- size\_t size ()
- std::vector< size\_t >::iterator begin ()
- std::vector< size\_t >::iterator end ()
- bool empty ()
- std::vector< double > \* get\_counters ()

## 8.34.1 Detailed Description

Definition at line 42 of file geese-types.hpp.

### 8.34.2 Constructor & Destructor Documentation

### 8.34.2.1 PhyloCounterData() [1/2]

Definition at line 48 of file geese-types.hpp.

### 8.34.2.2 PhyloCounterData() [2/2]

```
PhyloCounterData::PhyloCounterData ( ) [inline]
```

Definition at line 53 of file geese-types.hpp.

## 8.34.3 Member Function Documentation

#### 8.34.3.1 at()

Definition at line 55 of file geese-types.hpp.

## 8.34.3.2 begin()

```
std::vector< size_t >::iterator PhyloCounterData::begin ( ) [inline]
```

Definition at line 63 of file geese-types.hpp.

### 8.34.3.3 empty()

```
bool PhyloCounterData::empty ( ) [inline]
```

Definition at line 66 of file geese-types.hpp.

### 8.34.3.4 end()

```
std::vector< size_t >::iterator PhyloCounterData::end ( ) [inline]
```

Definition at line 64 of file geese-types.hpp.

### 8.34.3.5 get\_counters()

```
std::vector< double >* PhyloCounterData::get_counters ( ) [inline]
```

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

#### 8.34.3.6 operator()()

Definition at line 56 of file geese-types.hpp.

### 8.34.3.7 operator[]()

Definition at line 57 of file geese-types.hpp.

## 8.34.3.8 push\_back()

Definition at line 59 of file geese-types.hpp.

### 8.34.3.9 reserve()

Definition at line 58 of file geese-types.hpp.

#### 8.34.3.10 shrink\_to\_fit()

```
void PhyloCounterData::shrink_to_fit ( ) [inline]
```

Definition at line 60 of file geese-types.hpp.

### 8.34.3.11 size()

```
size_t PhyloCounterData::size ( ) [inline]
```

Definition at line 61 of file geese-types.hpp.

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

• include/barry/models/geese/geese-types.hpp

# 8.35 PhyloRuleDynData Class Reference

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

### **Public Member Functions**

- PhyloRuleDynData (const std::vector< double > \*counts\_, size\_t pos\_, size\_t lb\_, size\_t ub\_, size\_← t duplication\_)
- const double operator() () const
- ∼PhyloRuleDynData ()

### **Public Attributes**

- const std::vector< double > \* counts
- size\_t pos
- size\_t lb
- size\_t ub
- · size\_t duplication

# 8.35.1 Detailed Description

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

#### 8.35.2 Constructor & Destructor Documentation

## 8.35.2.1 PhyloRuleDynData()

Definition at line 79 of file geese-types.hpp.

## 8.35.2.2 ~PhyloRuleDynData()

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

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

## 8.35.3 Member Function Documentation

# 8.35.3.1 operator()()

```
const double PhyloRuleDynData::operator() ( ) const [inline]
```

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

## 8.35.4 Member Data Documentation

## 8.35.4.1 counts

```
const std::vector< double >* PhyloRuleDynData::counts
```

Definition at line 73 of file geese-types.hpp.

## 8.35.4.2 duplication

size\_t PhyloRuleDynData::duplication

Definition at line 77 of file geese-types.hpp.

#### 8.35.4.3 lb

size\_t PhyloRuleDynData::lb

Definition at line 75 of file geese-types.hpp.

## 8.35.4.4 pos

size\_t PhyloRuleDynData::pos

Definition at line 74 of file geese-types.hpp.

#### 8.35.4.5 ub

size\_t PhyloRuleDynData::ub

Definition at line 76 of file geese-types.hpp.

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

include/barry/models/geese/geese-types.hpp

# 8.36 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 (size\_t N\_, size\_t M\_)

#### Construct and destroy a PowerSet object

- PowerSet ()
- PowerSet (size\_t N\_, size\_t 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\_fun\_type < Array\_Type, Data\_Rule\_Type > count\_fun\_, Data\_Rule\_Type data\_)

## **Getter functions**

- const std::vector< Array\_Type > \* get\_data\_ptr () const
- std::vector< Array\_Type > get\_data () const
- std::vector< Array\_Type >::iterator begin ()
- std::vector< Array\_Type >::iterator end ()
- std::size\_t size () const noexcept
- const Array\_Type & operator[] (const size\_t &i) const

# **Public Attributes**

- Array\_Type EmptyArray
- $std::vector < Array\_Type > data$
- Rules
   Array\_Type, Data\_Rule\_Type > \* rules
- size\_t N
- size\_t M
- bool rules deleted = false
- std::vector < size\_t > coordinates\_free
- std::vector< size\_t > coordinates\_locked
- size\_t n\_free
- size\_t n\_locked

# 8.36.1 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 11 of file powerset-bones.hpp.

## 8.36.2 Constructor & Destructor Documentation

## 8.36.2.1 PowerSet() [1/3]

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

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

## 8.36.2.2 PowerSet() [2/3]

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

# 8.36.2.3 PowerSet() [3/3]

Definition at line 5 of file powerset-meat.hpp.

#### 8.36.2.4 ∼PowerSet()

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

Definition at line 13 of file powerset-meat.hpp.

# 8.36.3 Member Function Documentation

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

Definition at line 173 of file powerset-meat.hpp.

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

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

#### 8.36.3.3 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 68 of file powerset-bones.hpp.

# 8.36.3.4 calc()

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

Definition at line 144 of file powerset-meat.hpp.

#### 8.36.3.5 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 69 of file powerset-bones.hpp.

## 8.36.3.6 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 67 of file powerset-bones.hpp.

#### 8.36.3.7 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 66 of file powerset-bones.hpp.

#### 8.36.3.8 init support()

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

Definition at line 19 of file powerset-meat.hpp.

#### 8.36.3.9 operator[]()

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

#### 8.36.3.10 reset()

Definition at line 160 of file powerset-meat.hpp.

#### 8.36.3.11 size()

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

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

#### 8.36.4 Member Data Documentation

## 8.36.4.1 coordinates\_free

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

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

#### 8.36.4.2 coordinates locked

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

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

#### 8.36.4.3 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 19 of file powerset-bones.hpp.

#### 8.36.4.4 EmptyArray

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

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

#### 8.36.4.5 M

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

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

#### 8.36.4.6 N

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

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

# 8.36.4.7 n\_free

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

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

# 8.36.4.8 n\_locked

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

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

## 8.36.4.9 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 20 of file powerset-bones.hpp.

## 8.36.4.10 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 23 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

# 8.37 Progress Class Reference

A simple progress bar.

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

## **Public Member Functions**

```
• Progress (int n_, int width_)
```

- ∼Progress ()
- void next ()
- void end ()

# 8.37.1 Detailed Description

A simple progress bar.

Definition at line 11 of file progress.hpp.

## 8.37.2 Constructor & Destructor Documentation

# 8.37.2.1 Progress()

```
Progress::Progress (
          int n_,
          int width_ ) [inline]
```

Definition at line 30 of file progress.hpp.

## 8.37.2.2 ∼Progress()

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

Definition at line 23 of file progress.hpp.

# 8.37.3 Member Function Documentation

#### 8.37.3.1 end()

```
void Progress::end ( ) [inline]
```

Definition at line 52 of file progress.hpp.

# 8.37.3.2 next()

```
void Progress::next ( ) [inline]
```

Definition at line 41 of file progress.hpp.

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

• include/barry/progress.hpp

# 8.38 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 ()
- Data\_Type & D ()

Read/Write access to the data.

- bool operator() (const Array\_Type &a, size\_t i, size\_t j)
- std::string & get\_name ()
- std::string & get\_description ()
- std::string get\_name () const
- std::string get\_description () const

# 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\_, std::string name\_="", std::string desc ="")

# 8.38.1 Detailed Description

```
\label{template} $$ \ensuremath{\sf template}$$ < typename \ensuremath{\sf Array\_Type}$ = BArray<>, typename \ensuremath{\sf Data\_Type}$ = bool> class \ensuremath{\sf Rule}< Array\_Type, \ensuremath{\sf 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 20 of file rules-bones.hpp.

# 8.38.2 Constructor & Destructor Documentation

#### 8.38.2.1 Rule() [1/2]

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

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

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

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

## 8.38.2.3 ∼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.

#### 8.38.3 Member Function Documentation

# 8.38.3.1 D()

```
template<typename Array_Type , typename Data_Type >
Data_Type & Rule< Array_Type, Data_Type >::D [inline]
```

Read/Write access to the data.

Definition at line 37 of file rules-meat.hpp.

## 8.38.3.2 get\_description() [1/2]

```
template<typename Array_Type , typename Data_Type >
std::string & Rule< Array_Type, Data_Type >::get_description [inline]
```

Definition at line 54 of file rules-meat.hpp.

# 8.38.3.3 get\_description() [2/2]

```
template<typename Array_Type , typename Data_Type >
std::string Rule< Array_Type, Data_Type >::get_description [inline]
```

Definition at line 66 of file rules-meat.hpp.

# 8.38.3.4 get\_name() [1/2]

```
template<typename Array_Type , typename Data_Type >
std::string & Rule< Array_Type, Data_Type >::get_name [inline]
```

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

#### 8.38.3.5 get\_name() [2/2]

```
template<typename Array_Type , typename Data_Type >
std::string Rule< Array_Type, Data_Type >::get_name [inline]
```

Definition at line 60 of file rules-meat.hpp.

#### 8.38.3.6 operator()()

Definition at line 43 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

# 8.39 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 ()
- size\_t size () const noexcept
- bool operator() (const Array\_Type &a, size\_t i, size\_t j)

Check whether a given cell is free or locked.

- void get\_seq (const Array\_Type &a, std::vector< size\_t > \*free, std::vector< size\_t > \*locked=nullptr)

  Computes the sequence of free and locked cells in an BArray.
- std::vector< std::string > get\_names () const
- std::vector< std::string > get\_descriptions () const
- std::vector< Rule< Array\_Type, Data\_Type > >::iterator begin ()
- std::vector< Rule< Array\_Type, Data\_Type > >::iterator end ()

#### Rule adding

#### **Parameters**

```
rule
```

- void add rule (Rule < Array Type, Data Type > rule)
- void add\_rule (Rule\_fun\_type< Array\_Type, Data\_Type > rule\_, Data\_Type data\_, std::string name\_="", std::string description\_="")

# 8.39.1 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 71 of file rules-bones.hpp.

# 8.39.2 Constructor & Destructor Documentation

# 8.39.2.1 Rules() [1/2]

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

Definition at line 77 of file rules-bones.hpp.

## 8.39.2.2 Rules() [2/2]

Definition at line 5 of file rules-meat.hpp.

## 8.39.2.3 ∼Rules()

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

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

#### 8.39.3 Member Function Documentation

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

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

#### 8.39.3.2 add rule() [2/2]

Definition at line 82 of file rules-meat.hpp.

# 8.39.3.3 begin()

```
template<typename Array_Type , typename Data_Type >
std::vector< Rule<Array_Type,Data_Type> >::iterator Rules< Array_Type, Data_Type >::begin (
) [inline]
```

Definition at line 134 of file rules-bones.hpp.

# 8.39.3.4 end()

```
template<typename Array_Type , typename Data_Type >
std::vector< Rule<Array_Type,Data_Type> >::iterator Rules< Array_Type, Data_Type >::end ( )
[inline]
```

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

## 8.39.3.5 get\_descriptions()

```
template<typename Array_Type , typename Data_Type >
std::vector< std::string > Rules< Array_Type, Data_Type >::get_descriptions [inline]
```

Definition at line 179 of file rules-meat.hpp.

#### 8.39.3.6 get\_names()

```
template<typename Array_Type , typename Data_Type >
std::vector< std::string > Rules< Array_Type, Data_Type >::get_names [inline]
```

Definition at line 167 of file rules-meat.hpp.

# 8.39.3.7 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 117 of file rules-meat.hpp.

# 8.39.3.8 operator()()

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 101 of file rules-meat.hpp.

#### 8.39.3.9 operator=()

Definition at line 19 of file rules-meat.hpp.

# 8.39.3.10 size()

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

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

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

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

# 8.40 StatsCounter< Array\_Type, Data\_Type > Class Template Reference

Count stats for a single Array.

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

#### **Public Member Functions**

```
    StatsCounter (const Array_Type *Array_)
```

Creator of a StatsCounter

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

Copy constructor.

• 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 set\_counters (Counters < Array\_Type, Data\_Type > \*counters\_)
- void count\_init (size\_t i, size\_t 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 (size\_t i, size\_t j)
- std::vector< double > count\_all ()
- Counters < Array\_Type, Data\_Type > \* get\_counters ()
- std::vector< std::string > get\_names () const
- std::vector< std::string > get\_descriptions () const
- size t size () const

# 8.40.1 Detailed Description

```
\label{template} \begin{tabular}{ll} template < typename \ Array\_Type, \ typename \ Data\_Type > \\ class \ Stats Counter < Array\_Type, \ Data\_Type > \\ \end{tabular}
```

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 14 of file statscounter-bones.hpp.

#### 8.40.2 Constructor & Destructor Documentation

#### 8.40.2.1 StatsCounter() [1/3]

Creator of a StatsCounter

#### **Parameters**

Array←	A const pointer to a BArray.

Definition at line 37 of file statscounter-bones.hpp.

## 8.40.2.2 StatsCounter() [2/3]

Copy constructor.

#### **Parameters**

counter

# 8.40.2.3 StatsCounter() [3/3]

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

Can be created without setting the array.

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

# 8.40.2.4 $\sim$ StatsCounter()

```
template<typename Array_Type , typename Data_Type >
StatsCounter< Array_Type, Data_Type >::~StatsCounter ( )
```

# 8.40.3 Member Function Documentation

# 8.40.3.1 add\_counter()

## 8.40.3.2 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.

## 8.40.3.3 count\_current()

# 8.40.3.4 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.

# 8.40.3.5 get\_counters()

```
template<typename Array_Type , typename Data_Type >
Counters<Array_Type,Data_Type>* StatsCounter< Array_Type, Data_Type >::get_counters ( )
```

# 8.40.3.6 get\_descriptions()

```
template<typename Array_Type , typename Data_Type >
std::vector< std::string > StatsCounter< Array_Type, Data_Type >::get_descriptions ( ) const
```

#### 8.40.3.7 get\_names()

```
template<typename Array_Type , typename Data_Type >
std::vector< std::string > StatsCounter< Array_Type, Data_Type >::get_names ( ) const
```

# 8.40.3.8 reset\_array()

Changes the reference array for the counting.

#### **Parameters**

Array←	A pointer to an array of class Array_Type.

# 8.40.3.9 set\_counters()

#### 8.40.3.10 size()

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

Definition at line 86 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

# 8.41 Support < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data Rule Dyn Type > Class Template Reference

Compute the support of sufficient statistics.

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

#### **Public Member Functions**

Support (const Array\_Type &Array\_)

Constructor passing a reference Array.

• Support (size\_t N\_, size\_t M\_)

Constructor specifying the dimensions of the array (empty).

- Support ()
- ∼Support ()
- void init\_support (std::vector < Array\_Type > \*array\_bank=nullptr, std::vector < double > \*stats\_bank=nullptr)
- void calc (std::vector< Array\_Type > \*array\_bank=nullptr, std::vector< double > \*stats\_bank=nullptr, size
   \_t max\_num\_elements\_=0u)

Computes the entire support.

•  $std::vector < double > get\_counts$  () const

- std::vector< double > \* get\_current\_stats () List current statistics.
- void print () const
- const FregTable < double > & get\_data () const
- Counters < Array\_Type, Data\_Counter\_Type > \* get\_counters ()

Vector of couter functions.

Rules< Array\_Type, Data\_Rule\_Type > \* get\_rules ()

Vector of static rules (cells to iterate).

Rules< Array\_Type, Data\_Rule\_Dyn\_Type > \* get\_rules\_dyn ()

Vector of dynamic rules (to include/exclude a realizaton).

#### 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.

- $\bullet \ \ void \ add\_counter\ (Counter<Array\_Type,\ Data\_Counter\_Type>f\_)\\$
- 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\_)
- void add\_rule\_dyn (Rule< Array\_Type, Data\_Rule\_Dyn\_Type > \*f\_)
   void add\_rule\_dyn (Rule< Array\_Type, Data\_Rule\_Dyn\_Type > f\_)
- void set\_rules\_dyn (Rules < Array\_Type, Data\_Rule\_Dyn\_Type > \*rules\_)
- bool eval\_rules\_dyn (const std::vector< double > &counts, const size\_t &i, const size\_t &j)

#### **Public Attributes**

size\_t N

- size t M
- bool delete counters = true
- bool delete\_rules = true
- bool delete\_rules\_dyn = true
- size t max num elements = BARRY MAX NUM ELEMENTS
- std::vector< double > current stats
- std::vector< size t > coordinates free
- std::vector< size t > coordinates locked
- size\_t coordiantes\_n\_free
- size\_t coordiantes\_n\_locked
- std::vector< double > change stats
- std::vector< size\_t > hashes
- std::vector< bool > hashes initialized
- size\_t n\_counters

# 8.41.1 Detailed Description

```
template < typename Array_Type = BArray < bool, bool >, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool > class Support < Array_Type, Data_Counter_Type, Data_Rule_Dyn_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.

The members rule and rule\_dyn allow constraining the support. The first will establish which cells of the array will be used to iterate, for example, in the case of social networks, self-loops are not allowed, so the entire diagonal would be fixed to zero, reducing the size of the support.

In the case of  $rule_dyn$ , the function will stablish dynamically whether the current state will be included in the counts or not. For example, this set of rules can be used to constrain the support to networks that have a prescribed degree sequence.

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

# 8.41.2 Constructor & Destructor Documentation

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

Constructor passing a reference Array.

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

## 8.41.2.2 Support() [2/3]

Constructor specifying the dimensions of the array (empty).

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

# 8.41.2.3 Support() [3/3]

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

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

#### 8.41.2.4 ∼Support()

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

Definition at line 110 of file support-bones.hpp.

#### 8.41.3 Member Function Documentation

# 8.41.3.1 add\_counter()

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

#### 8.41.3.3 add rule() [2/2]

# 8.41.3.4 add\_rule\_dyn() [1/2]

# 8.41.3.5 add\_rule\_dyn() [2/2]

```
template<typename Array_Type = BArray<br/>bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool> void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::add_rule_\leftrightarrow dyn ( Rule< Array_Type, Data_Rule_Dyn_Type > f_)
```

## 8.41.3.6 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.

## 8.41.3.7 eval\_rules\_dyn()

# 8.41.3.8 get\_counters()

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename
Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Counters<Array_Type, Data_Counter_Type>* Support< Array_Type, Data_Counter_Type, Data_Rule_Dyn_Type >::get_counters ()
```

Vector of couter functions.

#### 8.41.3.9 get\_counts()

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< double > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn← _Type >::get_counts ( ) const
```

#### 8.41.3.10 get\_current\_stats()

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< double >* Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Cyn_Type >::get_current_stats ()
```

List current statistics.

## 8.41.3.11 get\_data()

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> const FreqTable< double >& Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_← Rule_Dyn_Type >::get_data ( ) const
```

## 8.41.3.12 get\_rules()

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename
Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Rules<Array_Type, Data_Rule_Type>* Support< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_rules ()
```

Vector of static rules (cells to iterate).

#### 8.41.3.13 get\_rules\_dyn()

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename
Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Rules<Array_Type, Data_Rule_Dyn_Type>* Support< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_rules_dyn ()
```

Vector of dynamic rules (to include/exclude a realizaton).

#### 8.41.3.14 init support()

# 8.41.3.15 print()

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename
Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::print ()
const
```

## 8.41.3.16 reset\_array() [1/2]

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename
Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::reset_array
( )
```

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

# 8.41.3.18 set\_counters()

# 8.41.3.19 set\_rules()

#### 8.41.3.20 set\_rules\_dyn()

# 8.41.4 Member Data Documentation

# 8.41.4.1 change\_stats

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< double > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn← _Type >::change_stats
```

Definition at line 80 of file support-bones.hpp.

#### 8.41.4.2 coordiantes\_n\_free

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::coordiantes ← __n_free
```

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

#### 8.41.4.3 coordiantes n locked

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::coordiantes 
n locked
```

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

#### 8.41.4.4 coordinates\_free

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< size_t > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn 
_Type >::coordinates_free
```

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

## 8.41.4.5 coordinates\_locked

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< size_t > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn← _Type >::coordinates_locked
```

Definition at line 77 of file support-bones.hpp.

#### 8.41.4.6 current stats

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< double > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn← _Type >::current_stats
```

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

#### 8.41.4.7 delete\_counters

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_← counters = true
```

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

#### 8.41.4.8 delete rules

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_← rules = true
```

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

#### 8.41.4.9 delete\_rules\_dyn

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_← rules_dyn = true
```

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

#### 8.41.4.10 hashes

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< size_t > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn← _Type >::hashes
```

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

#### 8.41.4.11 hashes initialized

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< bool > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_↔
Type >::hashes_initialized
```

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

#### 8.41.4.12 M

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename
Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::M
```

Definition at line 68 of file support-bones.hpp.

#### 8.41.4.13 max\_num\_elements

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::max_num - elements = BARRY_MAX_NUM_ELEMENTS
```

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

# 8.41.4.14 N

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename
Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::N
```

Definition at line 68 of file support-bones.hpp.

#### 8.41.4.15 n counters

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::n_
counters
```

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

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

include/barry/support-bones.hpp

# 8.42 vecHasher < T > Struct Template Reference

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

# **Public Member Functions**

• std::size\_t operator() (std::vector< T > const &dat) const noexcept

# 8.42.1 Detailed Description

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

Definition at line 105 of file typedefs.hpp.

# 8.42.2 Member Function Documentation

# 8.42.2.1 operator()()

Definition at line 108 of file typedefs.hpp.

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

• include/barry/typedefs.hpp

# **Chapter 9**

# **File Documentation**

# 9.1 include/barry/barray-bones.hpp File Reference

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



# **Classes**

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

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

## Classes

class ConstBArrayRowIter< Cell\_Type, Data\_Type >

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

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



## **Macros**

- #define BARRAY\_TYPE() BArray<Cell\_Type, Data\_Type>
- #define BARRAY\_TEMPLATE\_ARGS() < typename Cell\_Type, typename Data\_Type>
- #define BARRAY\_TEMPLATE(a, b) template BARRAY\_TEMPLATE\_ARGS() inline a BARRAY\_TYPE()::b
- #define ROW(a) this->el\_ij[a]
- #define COL(a) this->el\_ji[a]

#### **Functions**

- template BARRAY TEMPLATE ARGS () inline void checkdim (const BARRAY TYPE() &lhs
- template const BARRAY TYPE () &rhs)
- BARRAY\_TEMPLATE (BARRAY\_TYPE()&, operator+=)(const BArray< Cell\_Type
- for (size\_t i=0u;i< nrow();++i) for(size\_t j=0u = el[POS(i, j)]
- j< ncol();++j) this-> operator() (i, j)+
- BARRAY\_TEMPLATE (BARRAY\_TYPE()&, operator+=)(const Cell\_Type &rhs)
- BARRAY\_TEMPLATE (BARRAY\_TYPE()&, operator-=)(const BArray< Cell\_Type
- BARRAY\_TEMPLATE (BARRAY\_TYPE()&, operator-=)(const Cell\_Type &rhs)
- BARRAY\_TEMPLATE (BARRAY\_TYPE()&, operator\*=)(const Cell\_Type &rhs)
- BARRAY\_TEMPLATE (BARRAY\_TYPE()&, operator/=)(const Cell\_Type &rhs)

# **Variables**

- · Data\_Type & rhs
- return \* this

#### 9.3.1 Macro Definition Documentation

# 9.3.1.1 BARRAY\_TEMPLATE

Definition at line 11 of file barray-meat-operators.hpp.

# 9.3.1.2 BARRAY\_TEMPLATE\_ARGS

```
template BARRAY_TEMPLATE_ARGS() <typename Cell_Type, typename Data_Type>
```

Definition at line 9 of file barray-meat-operators.hpp.

# 9.3.1.3 BARRAY\_TYPE

```
template Data_Type BARRAY_TYPE( ) BArray<Cell_Type, Data_Type>
```

Definition at line 7 of file barray-meat-operators.hpp.

# 9.3.1.4 COL

Definition at line 15 of file barray-meat-operators.hpp.

#### 9.3.1.5 ROW

Definition at line 14 of file barray-meat-operators.hpp.

# 9.3.2 Function Documentation

244 File Documentation

# 9.3.2.1 BARRAY\_TEMPLATE() [1/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator* ) const &
```

Definition at line 88 of file barray-meat-operators.hpp.

# 9.3.2.2 BARRAY\_TEMPLATE() [2/6]

# 9.3.2.3 BARRAY\_TEMPLATE() [3/6]

Definition at line 46 of file barray-meat-operators.hpp.

# 9.3.2.4 BARRAY\_TEMPLATE() [4/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator- ) const
```

# 9.3.2.5 BARRAY\_TEMPLATE() [5/6]

Definition at line 75 of file barray-meat-operators.hpp.

# **9.3.2.6 BARRAY\_TEMPLATE()** [6/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator/ ) const &
```

Definition at line 105 of file barray-meat-operators.hpp.

### 9.3.2.7 BARRAY\_TEMPLATE\_ARGS()

```
template BARRAY_TEMPLATE_ARGS ( ) const \&
```

### 9.3.2.8 BARRAY\_TYPE()

```
template const BARRAY_TYPE ( ) &
```

Definition at line 20 of file barray-meat-operators.hpp.

#### 9.3.2.9 for()

```
for ( ) = el[POS(i, j)] [pure virtual]
```

Definition at line 66 of file barray-meat-operators.hpp.

### 9.3.2.10 operator()()

# 9.3.3 Variable Documentation

#### 9.3.3.1 rhs

```
Data_Type & rhs
Initial value:
{
    checkdim_(*this, rhs)
```

Definition at line 33 of file barray-meat-operators.hpp.

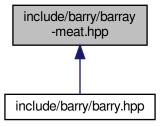
#### 9.3.3.2 this

```
return * this
```

Definition at line 43 of file barray-meat-operators.hpp.

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

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



### **Macros**

- #define BARRAY\_TYPE() BArray<Cell\_Type, Data\_Type>
- #define BARRAY\_TEMPLATE\_ARGS() < typename Cell\_Type, typename Data\_Type>
- #define BARRAY\_TEMPLATE(a, b) template BARRAY\_TEMPLATE\_ARGS() inline a BARRAY\_TYPE()::b
- #define ROW(a) this->el\_ij[a]
- #define COL(a) this->el\_ji[a]

#### **Functions**

```
• BARRAY TEMPLATE (, BArray)(size_t N_
• el ij resize (N)
• el_ji resize (M)

    for (size t i=0u;i < source.size();++i)</li>

    Data Type bool M (Array .M)

    BARRAY_TEMPLATE (BARRAY_TYPE() &, operator=)(const BArray< Cell_Type</li>

    BARRAY_TEMPLATE (, BArray)(BARRAY_TYPE() &&x) noexcept

    BARRAY TEMPLATE (BARRAY TYPE() &, operator=)(BARRAY TYPE() &&x) noexcept

• BARRAY TEMPLATE (bool, operator==)(const BARRAY TYPE() & Array )

    BARRAY TEMPLATE (,~BArray)()

    BARRAY_TEMPLATE (void, set_data)(Data_Type *data_

    BARRAY TEMPLATE (Data Type *, D ptr)()

• BARRAY_TEMPLATE (Data_Type &, D)()

    BARRAY TEMPLATE (void, out of range)(size ti

    BARRAY TEMPLATE (Cell Type, get cell)(size ti

    if (ROW(i).size()==0u) return(Cell_Type) 0.0

• if (search !=ROW(i).end()) return search -> second.value
• return (Cell_Type) 0.0

    BARRAY_TEMPLATE (std::vector< Cell_Type >, get_row_vec)(size_t i

    std::vector< Cell Type > ans (ncol(),(Cell Type) false)

    for (const auto &iter :row(i, false)) ans[iter.first]

    BARRAY_TEMPLATE (void, get_row_vec)(std

    BARRAY_TEMPLATE (BARRAY_TYPE() &, operator-=)(const std

• BARRAY_TEMPLATE (void, insert_cell)(size_t i
· if (check exists)

    COL (j).emplace(i

• & ROW (i)[j])

    BARRAY_TEMPLATE (void, swap_cells)(size_t i0

if (report !=nullptr)(*report)
• if (check0 &check1)

    else if (!check0 &check1)

• else if (check0 &!check1)

    BARRAY TEMPLATE (void, toggle cell)(size ti

• BARRAY_TEMPLATE (void, swap_rows)(size_t i0
• if (ROW(i0).size()==0u) move0
• if (ROW(i1).size()==0u) move1
• if (!move0 &&!move1) return

    ROW (i0).swap(ROW(i1))

    BARRAY_TEMPLATE (void, swap_cols)(size_t j0

• if (COL(j0).size()==0u) check0
if (COL(j1).size()==0u) check1

    if (check0 &&check1)

    else if (check0 &&!check1)

    else if (!check0 &&check1)

• BARRAY_TEMPLATE (void, zero_row)(size_t i

    for (auto row=row0.begin();row !=row0.end();++row) rm cell(i

    BARRAY_TEMPLATE (void, zero_col)(size_t j

    if (COL(j).size()==0u) return

• BARRAY_TEMPLATE (void, transpose)()

    BARRAY_TEMPLATE (void, clear)(bool hard)

    BARRAY_TEMPLATE (void, resize)(size_t N_

    if (M_< M) for(size_t j = N_</li>
```

#### **Variables**

```
 size t M

    size_t const std::vector < size_t > & source

    size_t const std::vector< size_t > const std::vector< size_t > & target

 \bullet \  \, \text{size\_t const std::vector} < \  \, \text{size\_t} > \  \, \text{const std::vector} < \  \, \text{cize\_t} > \  \, \text{const std::vector} < \  \, \text{Cell\_Type} > \  \, \& \  \, \text{value} 
• size t const std::vector< size t > const std::vector< size t > const std::vector< Cell Type > bool add
• if(source.size() !=value.size()) throw std N = N_

    M = M

    return

    Data_Type & Array_

    Data_Type bool copy_data

• bool delete_data_

 data = data

    delete data = delete data

• size t j const
size_t j
• auto search = ROW(i).find(j)

    return ans

    size_t const Cell
    Cell_Type > & v

    size_t const Cell< Cell_Type > bool check_bounds

    size_t const Cell
    Cell_Type > bool bool check_exists

    else

    NCells

• size_t j0
size_t size_t i1
• size_t size_t j1
• size_t size_t bool int int * report
• auto row0 = ROW(i)

    row first

    row false

auto col0 = COL(j)
```

#### 9.4.1 Macro Definition Documentation

#### 9.4.1.1 BARRAY\_TEMPLATE

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

#### 9.4.1.2 BARRAY\_TEMPLATE\_ARGS

```
#define BARRAY_TEMPLATE_ARGS( ) <typename Cell_Type, typename Data_Type>
```

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

# 9.4.1.3 BARRAY\_TYPE

```
#define BARRAY_TYPE( ) BArray<Cell_Type, Data_Type>
```

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

#### 9.4.1.4 COL

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

#### 9.4.1.5 ROW

Definition at line 20 of file barray-meat.hpp.

### 9.4.2 Function Documentation

### 9.4.2.1 ans()

# 9.4.2.2 BARRAY\_TEMPLATE() [1/24]

```
BARRAY_TEMPLATE (

BArray ) && [noexcept]
```

Definition at line 230 of file barray-meat.hpp.

# 9.4.2.3 BARRAY\_TEMPLATE() [2/24]

```
BARRAY_TEMPLATE (
BArray )
```

### 9.4.2.4 BARRAY\_TEMPLATE() [3/24]

```
BARRAY_TEMPLATE ( \sim \textit{BArray} \ )
```

Definition at line 339 of file barray-meat.hpp.

# 9.4.2.5 BARRAY\_TEMPLATE() [4/24]

Definition at line 597 of file barray-meat.hpp.

# 9.4.2.6 BARRAY\_TEMPLATE() [5/24]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE() & ,
          operator ) && [noexcept]
```

Definition at line 272 of file barray-meat.hpp.

#### 9.4.2.7 BARRAY\_TEMPLATE() [6/24]

### 9.4.2.8 BARRAY\_TEMPLATE() [7/24]

```
BARRAY_TEMPLATE (
          bool ,
          operator = = ) const &
```

Definition at line 321 of file barray-meat.hpp.

# 9.4.2.9 BARRAY\_TEMPLATE() [8/24]

# 9.4.2.10 BARRAY\_TEMPLATE() [9/24]

```
BARRAY_TEMPLATE (

Data_Type & ,

D )
```

Definition at line 372 of file barray-meat.hpp.

### 9.4.2.11 BARRAY\_TEMPLATE() [10/24]

Definition at line 361 of file barray-meat.hpp.

### 9.4.2.12 BARRAY\_TEMPLATE() [11/24]

```
BARRAY_TEMPLATE (
          std::vector< Cell_Type > ,
          get_row_vec )
```

### 9.4.2.13 BARRAY\_TEMPLATE() [12/24]

```
BARRAY_TEMPLATE ( void , clear )
```

Definition at line 1130 of file barray-meat.hpp.

# 9.4.2.14 BARRAY\_TEMPLATE() [13/24]

Definition at line 452 of file barray-meat.hpp.

# 9.4.2.15 BARRAY\_TEMPLATE() [14/24]

### 9.4.2.16 BARRAY\_TEMPLATE() [15/24]

# 9.4.2.17 BARRAY\_TEMPLATE() [16/24]

# 9.4.2.18 BARRAY\_TEMPLATE() [17/24]

### 9.4.2.19 BARRAY\_TEMPLATE() [18/24]

# 9.4.2.20 BARRAY\_TEMPLATE() [19/24]

# 9.4.2.21 BARRAY\_TEMPLATE() [20/24]

# 9.4.2.22 BARRAY\_TEMPLATE() [21/24]

#### 9.4.2.23 BARRAY\_TEMPLATE() [22/24]

Definition at line 1069 of file barray-meat.hpp.

# 9.4.2.24 BARRAY\_TEMPLATE() [23/24]

```
BARRAY_TEMPLATE ( void , zero_col )
```

### 9.4.2.25 BARRAY\_TEMPLATE() [24/24]

```
BARRAY_TEMPLATE (

void ,

zero_row )
```

```
9.4.2.26 COL()
```

```
COL (
```

# **9.4.2.27** for() [1/3]

```
for (
    auto row = row0.begin();row !=row0.end();++row )
```

#### 9.4.2.28 for() [2/3]

### 9.4.2.29 for() [3/3]

```
for ( )
```

Definition at line 51 of file barray-meat.hpp.

#### 9.4.2.30 if() [1/17]

```
else if ( !check0 && check1 )
```

Definition at line 1008 of file barray-meat.hpp.

# 9.4.2.31 if() [2/17]

```
else if (
    !check0 & check1 )
```

Definition at line 856 of file barray-meat.hpp.

# 9.4.2.32 if() [3/17]

```
if (
    !move0 &&! move1 )
```

### 9.4.2.33 if() [4/17]

Definition at line 864 of file barray-meat.hpp.

# 9.4.2.34 if() [5/17]

```
else if (
          check0 &&! check1 )
```

Definition at line 999 of file barray-meat.hpp.

#### 9.4.2.35 if() [6/17]

```
if ( check0 && check1)
```

Definition at line 972 of file barray-meat.hpp.

### 9.4.2.36 if() [7/17]

```
if ( check0 & check1)
```

Definition at line 838 of file barray-meat.hpp.

### 9.4.2.37 if() [8/17]

```
else if (
          check_exists = = CHECK::BOTH )
```

Definition at line 679 of file barray-meat.hpp.

```
9.4.2.38 if() [9/17]
```

```
if ( COL(j).size() = =0u )
```

# **9.4.2.39** if() [10/17]

```
if ( \label{eq:col} \texttt{COL(j0).size()} \quad = = 0u \ )
```

# 9.4.2.40 if() [11/17]

```
if ( COL(j1).size() = =0u )
```

#### 9.4.2.41 if() [12/17]

```
else if ( ) = N_
```

Definition at line 86 of file barray-meat.hpp.

# 9.4.2.42 if() [13/17]

```
if (  {\tt report !} \quad = {\tt nullptr} \; ) \\
```

# 9.4.2.43 if() [14/17]

```
if ( \label{eq:row_row_row} \mbox{ROW(i).size()} = = 0u \mbox{ )}
```

# **9.4.2.44** if() [15/17]

```
if ( \label{eq:row_row_row} \mbox{ROW(iO).size()} \ = \ = \mbox{$0$$u$} \ )
```

# 9.4.2.45 if() [16/17]

```
if ( \label{eq:row_row_row} \mbox{ROW(i1).size()} = = 0 \mbox{$u$} \mbox{ )}
```

# 9.4.2.46 if() [17/17]

```
if (
    search ! = ROW(i).end() ) -> second.value
```

# 9.4.2.47 M()

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

# 9.4.2.48 resize() [1/2]

```
el_ji resize (
M )
```

### 9.4.2.49 resize() [2/2]

```
el_ij resize (
N )
```

### 9.4.2.50 return()

# 9.4.2.51 ROW() [1/2]

```
& ROW ( i )
```

### 9.4.2.52 ROW() [2/2]

```
ROW ( i0 )
```

### 9.4.3 Variable Documentation

#### 9.4.3.1 add

```
size_t const std::vector< size_t > const std::vector< size_t > bool add

Initial value:
{
    if (source.size() != target.size())
        throw std::length_error("-source- and -target- don't match on length.")
```

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

#### 9.4.3.2 ans

return ans

Definition at line 449 of file barray-meat.hpp.

#### 9.4.3.3 Array\_

Data\_Type & Array\_

Definition at line 134 of file barray-meat.hpp.

# 9.4.3.4 check\_bounds

```
bool check_bounds

Initial value:
{
    if (check_bounds) {
        out_of_range(i0,0u);
        out_of_range(i1,0u);
    }
    bool move0=true, move1=true
```

Definition at line 672 of file barray-meat.hpp.

## 9.4.3.5 check\_exists

```
size_t bool int check_exists

Initial value:
{
    if (check_bounds)
        out_of_range(i,j)
```

Definition at line 673 of file barray-meat.hpp.

### 9.4.3.6 col0

```
auto col0 = COL(j)
```

Definition at line 1061 of file barray-meat.hpp.

#### 9.4.3.7 const

```
size_t bool check_bounds const

Initial value:
{
    if (i >= N)
        throw std::range_error("The row is out of range.")
```

Definition at line 402 of file barray-meat.hpp.

# 9.4.3.8 copy\_data

```
Data_Type bool copy_data
```

Definition at line 135 of file barray-meat.hpp.

### 9.4.3.9 data

```
data = data_
```

Definition at line 354 of file barray-meat.hpp.

#### 9.4.3.10 delete\_data

```
delete_data = delete_data_
```

Definition at line 355 of file barray-meat.hpp.

# 9.4.3.11 delete\_data\_

Definition at line 348 of file barray-meat.hpp.

#### 9.4.3.12 else

Definition at line 703 of file barray-meat.hpp.

#### 9.4.3.13 false

row false

Definition at line 1042 of file barray-meat.hpp.

### 9.4.3.14 first

```
row first
```

Definition at line 1042 of file barray-meat.hpp.

#### 9.4.3.15 i1

```
size_t i1
```

Definition at line 776 of file barray-meat.hpp.

#### 9.4.3.16 j

```
size_t j
```

#### Initial value:

```
if (init_fun == nullptr)
    return 0.0
```

Definition at line 414 of file barray-meat.hpp.

#### 9.4.3.17 j0

```
size_t j0
```

Definition at line 775 of file barray-meat.hpp.

## 9.4.3.18 j1

```
size_t j1
```

Definition at line 776 of file barray-meat.hpp.

#### 9.4.3.19 M

```
M = M_{\underline{}}
```

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

### 9.4.3.20 M\_

```
size_t M_
```

#### Initial value:

ililiai va

```
if (N_ < N)
    for (size_t i = N_; i < N; ++i)
        zero_row(i, false)</pre>
```

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

#### 9.4.3.21 N

```
if (source.size() != target.size()) throw std if (source.size() != value.size()) throw std N = N
```

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

#### 9.4.3.22 NCells

NCells

Definition at line 707 of file barray-meat.hpp.

#### 9.4.3.23 report

```
size_t size_t size_t bool int int* report
```

Definition at line 779 of file barray-meat.hpp.

#### 9.4.3.24 return

return

Definition at line 66 of file barray-meat.hpp.

# 9.4.3.25 row0

```
auto row0 = ROW(i)
```

Definition at line 1040 of file barray-meat.hpp.

#### 9.4.3.26 search

```
auto search = ROW(i).find(j)
```

Definition at line 426 of file barray-meat.hpp.

#### 9.4.3.27 source

```
size_t const std::vector< size_t > & source
```

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

## 9.4.3.28 target

```
size_t const std::vector< size_t > const std::vector< size_t > & target
```

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

#### 9.4.3.29 v

```
size_t Cell_Type v
```

Definition at line 671 of file barray-meat.hpp.

#### 9.4.3.30 value

size\_t const std::vector< size\_t > const std::vector< Size\_t > const std::vector< Cell\_Type
>& value

Definition at line 33 of file barray-meat.hpp.

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

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 >

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

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



# 9.7 include/barry/barraydense-bones.hpp File Reference

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



#### Classes

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

# 9.8 include/barry/barraydense-meat-operators.hpp File Reference

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



#### **Macros**

- #define BDENSE\_TYPE() BArrayDense<Cell\_Type, Data\_Type>
- #define BDENSE\_TEMPLATE\_ARGS() < typename Cell\_Type, typename Data\_Type>
- #define BDENSE\_TEMPLATE(a, b) template BDENSE\_TEMPLATE\_ARGS() inline a BDENSE\_TYPE()::b
- #define ROW(a) this->el\_ij[a]
- #define COL(a) this->el\_ji[a]
- #define POS(a, b) (b)\*N + (a)
- #define POS\_N(a, b, c) (b)\*(c) + (a)

#### **Functions**

- template BDENSE\_TEMPLATE\_ARGS () inline void checkdim\_(const BDENSE\_TYPE() &lhs
- template const BDENSE\_TYPE () &rhs)
- BDENSE\_TEMPLATE (BDENSE\_TYPE()&, operator+=)(const BDENSE\_TYPE() &rhs)
- BDENSE TEMPLATE (BDENSE TYPE()&, operator-=)(const BDENSE TYPE() &rhs)
- BDENSE\_TEMPLATE (BDENSE\_TYPE()&, operator\*=)(const Cell\_Type &rhs)
- BDENSE\_TEMPLATE (BDENSE\_TYPE()&, operator/=)(const Cell\_Type &rhs)

#### 9.8.1 Macro Definition Documentation

#### 9.8.1.1 BDENSE TEMPLATE

Definition at line 11 of file barraydense-meat-operators.hpp.

#### 9.8.1.2 BDENSE\_TEMPLATE\_ARGS

```
template BDENSE_TEMPLATE_ARGS() <typename Cell_Type, typename Data_Type>
```

Definition at line 9 of file barraydense-meat-operators.hpp.

#### 9.8.1.3 BDENSE\_TYPE

```
template Data_Type BDENSE_TYPE( ) BArrayDense<Cell_Type, Data_Type>
```

Definition at line 7 of file barraydense-meat-operators.hpp.

### 9.8.1.4 COL

Definition at line 15 of file barraydense-meat-operators.hpp.

#### 9.8.1.5 POS

```
#define POS(  a, \\ b ) \ (b)*N + (a)
```

Definition at line 16 of file barraydense-meat-operators.hpp.

### 9.8.1.6 POS\_N

Definition at line 17 of file barraydense-meat-operators.hpp.

#### 9.8.1.7 ROW

Definition at line 14 of file barraydense-meat-operators.hpp.

#### 9.8.2 Function Documentation

### 9.8.2.1 BDENSE\_TEMPLATE() [1/4]

Definition at line 90 of file barraydense-meat-operators.hpp.

### 9.8.2.2 BDENSE\_TEMPLATE() [2/4]

Definition at line 34 of file barraydense-meat-operators.hpp.

### 9.8.2.3 BDENSE\_TEMPLATE() [3/4]

Definition at line 61 of file barraydense-meat-operators.hpp.

## 9.8.2.4 BDENSE\_TEMPLATE() [4/4]

Definition at line 101 of file barraydense-meat-operators.hpp.

#### 9.8.2.5 BDENSE\_TEMPLATE\_ARGS()

```
template BDENSE_TEMPLATE_ARGS ( ) const \&
```

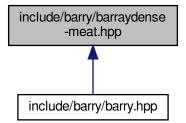
#### 9.8.2.6 BDENSE\_TYPE()

```
template const BDENSE_TYPE ( ) &
```

Definition at line 22 of file barraydense-meat-operators.hpp.

# 9.9 include/barry/barraydense-meat.hpp File Reference

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



#### Macros

```
    #define BDENSE_TYPE() BArrayDense<Cell_Type, Data_Type>
```

- #define BDENSE\_TEMPLATE\_ARGS() < typename Cell\_Type, typename Data\_Type>
- #define BDENSE\_TEMPLATE(a, b) template BDENSE\_TEMPLATE\_ARGS() inline a BDENSE\_TYPE()::b
- #define ROW(a) this->el\_ij[a]
- #define COL(a) this->el ji[a]
- #define POS(a, b) (b)\*N + (a)
- #define POS\_N(a, b, c) (b)\*(c) + (a)
- #define ZERO\_CELL static\_cast<Cell\_Type>(0.0)

#### **Functions**

```
• BDENSE_TEMPLATE (, BArrayDense)(size_t N_
```

- el resize (N \*M, ZERO\_CELL)
- el rowsums resize (N, ZERO CELL)
- el\_colsums resize (M, ZERO\_CELL)
- for (size\_t i=0u;i < source.size();++i)</li>
- BDENSE\_TEMPLATE (, BArrayDense)(const BDENSE\_TYPE() &Array\_
- bool M (Array .M)
- BDENSE\_TEMPLATE (BDENSE\_TYPE() &, operator=)(const BDENSE\_TYPE() &Array\_)
- BDENSE TEMPLATE (, BArrayDense)(BDENSE TYPE() &&x) noexcept
- BDENSE\_TEMPLATE (BDENSE\_TYPE() &, operator=)(BDENSE\_TYPE() &&x) noexcept
- BDENSE\_TEMPLATE (bool, operator==)(const BDENSE\_TYPE() &Array\_)
- BDENSE TEMPLATE (, ~BArrayDense)()
- BDENSE\_TEMPLATE (void, set\_data)(Data\_Type \*data\_
- BDENSE\_TEMPLATE (Data\_Type \*, D\_ptr)()
- BDENSE TEMPLATE (const Data Type \*, D ptr)() const
- BDENSE TEMPLATE (Data Type &, D)()
- BDENSE\_TEMPLATE (const Data\_Type &, D)() const
- BDENSE\_TEMPLATE (void, out\_of\_range)(size\_t i
- BDENSE\_TEMPLATE (Cell\_Type, get\_cell)(size\_t i
- BDENSE\_TEMPLATE (std::vector< Cell\_Type >, get\_row\_vec)(size\_t i
- std::vector< Cell\_Type > ans (ncol(), static\_cast< Cell\_Type >(false))
- BDENSE\_TEMPLATE (void, get\_row\_vec)(std
- BDENSE\_TEMPLATE (Entries < Cell\_Type >, get\_entries)() const
- BDENSE\_TEMPLATE (bool, is\_empty)(size\_t i
- BDENSE\_TEMPLATE (size\_t, nrow)() const noexcept
- BDENSE\_TEMPLATE (size\_t, ncol)() const noexcept
- BDENSE\_TEMPLATE (size\_t, nnozero)() const noexcept
- BDENSE\_TEMPLATE (Cell< Cell\_Type >, default\_val)() const
- BDENSE\_TEMPLATE (BDENSE\_TYPE() &, operator+=)(const std
- BDENSE\_TEMPLATE (BDENSE\_TYPE() &, operator-=)(const std
- BDENSE\_TEMPLATE (void, insert\_cell)(size\_t i
- if (el[POS(i, j)]==BARRY\_ZERO\_DENSE)
- BDENSE\_TEMPLATE (void, swap\_cells)(size\_t i0
- if ((i0==i1) &&(j0==j1)) return
- rm cell (i0, j0, false, false)
- rm\_cell (i1, j1, false, false)
- insert\_cell (i0, j0, val1, false, false)
- insert\_cell (i1, j1, val0, false, false)
- BDENSE\_TEMPLATE (void, toggle\_cell)(size\_t i
- else rm\_cell (i, j, false, false)
- BDENSE\_TEMPLATE (void, swap\_rows)(size\_t i0

```
• BDENSE_TEMPLATE (void, swap_cols)(size_t j0

    BDENSE_TEMPLATE (void, zero_row)(size_t i

• if (el_rowsums[i]==ZERO_CELL) return
• BDENSE_TEMPLATE (void, zero_col)(size_t j
• if (el colsums[j]==ZERO CELL) return
• BDENSE_TEMPLATE (void, transpose)()
• BDENSE_TEMPLATE (void, clear)(bool hard)

    BDENSE_TEMPLATE (void, resize)(size_t N_

• el resize (N_ *M_, ZERO_CELL)
• el rowsums resize (N, ZERO CELL)
• el_colsums resize (M_, ZERO_CELL)
• BDENSE TEMPLATE (void, reserve)()

    BDENSE_TEMPLATE (void, print)(const char *fmt

• va_start (args, fmt)
• printf barry (fmt, args)

    va end (args)

• BDENSE TEMPLATE (const std::vector< Cell Type > &, get data)() const

    BDENSE_TEMPLATE (const Cell_Type, rowsum)(size_t i) const

• BDENSE_TEMPLATE (const Cell_Type, colsum)(size_t j) const
```

#### **Variables**

col

```
 size t M

• size t const std::vector< size t > & source

    size_t const std::vector< size_t > const std::vector< size_t > & target

• size_t const std::vector< size_t > const std::vector< Cell_Type > & value

    size_t const std::vector < size_t > const std::vector < cell_Type > bool add

• if(source.size() !=value.size()) throw std N = N_
• M = M

    return

· bool copy data

    bool delete_data_

data = data
• delete data = delete data
· size t i const
size_t j

    return el [POS(i, j)] == ZERO CELL

· return ans

    size_t const Cell
    Cell_Type > & v

    size_t const Cell
    Cell_Type > bool check_bounds

    size_t const Cell< Cell_Type > bool bool check_exists

    else

• el_rowsums [i] = (v.value - old)
el_colsums [j] = (v.value - old)
size_t j0
size_t size_t i1
• size t size t size t i1

    size_t size_t bool int int * report

    Cell_Type val0 = el[POS(i0,j0)]

Cell_Type val1 = el[POS(i1,j1)]
· false
```

#### 9.9.1 Macro Definition Documentation

### 9.9.1.1 BDENSE\_TEMPLATE

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

#### 9.9.1.2 BDENSE\_TEMPLATE\_ARGS

```
#define BDENSE_TEMPLATE_ARGS( ) <typename Cell_Type, typename Data_Type>
```

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

#### 9.9.1.3 BDENSE TYPE

```
#define BDENSE_TYPE( ) BArrayDense<Cell_Type, Data_Type>
```

Definition at line 23 of file barraydense-meat.hpp.

#### 9.9.1.4 COL

Definition at line 31 of file barraydense-meat.hpp.

### 9.9.1.5 POS

```
#define POS(  a, \\ b ) (b)*N + (a)
```

Definition at line 32 of file barraydense-meat.hpp.

### 9.9.1.6 POS\_N

Definition at line 33 of file barraydense-meat.hpp.

#### 9.9.1.7 ROW

Definition at line 30 of file barraydense-meat.hpp.

#### 9.9.1.8 ZERO CELL

```
#define ZERO_CELL static_cast<Cell_Type>(0.0)
```

Definition at line 38 of file barraydense-meat.hpp.

### 9.9.2 Function Documentation

#### 9.9.2.1 ans()

### **9.9.2.2 BDENSE\_TEMPLATE()** [1/39]

Definition at line 240 of file barraydense-meat.hpp.

### 9.9.2.3 BDENSE\_TEMPLATE() [2/39]

#### 9.9.2.4 BDENSE\_TEMPLATE() [3/39]

```
BDENSE_TEMPLATE (

BArrayDense )
```

#### 9.9.2.5 BDENSE\_TEMPLATE() [4/39]

```
BDENSE_TEMPLATE ( \sim \textit{BArrayDense} \ )
```

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

#### 9.9.2.6 BDENSE\_TEMPLATE() [5/39]

Definition at line 566 of file barraydense-meat.hpp.

#### 9.9.2.7 BDENSE\_TEMPLATE() [6/39]

Definition at line 584 of file barraydense-meat.hpp.

### 9.9.2.8 BDENSE\_TEMPLATE() [7/39]

Definition at line 257 of file barraydense-meat.hpp.

### 9.9.2.9 BDENSE\_TEMPLATE() [8/39]

Definition at line 194 of file barraydense-meat.hpp.

### **9.9.2.10 BDENSE\_TEMPLATE()** [9/39]

```
BDENSE_TEMPLATE (
          bool ,
          is_empty )
```

### **9.9.2.11 BDENSE\_TEMPLATE()** [10/39]

```
BDENSE_TEMPLATE (
          bool ,
          operator = = ) const &
```

Definition at line 300 of file barraydense-meat.hpp.

### 9.9.2.12 BDENSE\_TEMPLATE() [11/39]

Definition at line 562 of file barraydense-meat.hpp.

#### 9.9.2.13 BDENSE\_TEMPLATE() [12/39]

# **9.9.2.14 BDENSE\_TEMPLATE()** [13/39]

Definition at line 999 of file barraydense-meat.hpp.

### 9.9.2.15 BDENSE\_TEMPLATE() [14/39]

Definition at line 994 of file barraydense-meat.hpp.

#### 9.9.2.16 BDENSE\_TEMPLATE() [15/39]

Definition at line 353 of file barraydense-meat.hpp.

#### 9.9.2.17 BDENSE\_TEMPLATE() [16/39]

Definition at line 345 of file barraydense-meat.hpp.

### **9.9.2.18 BDENSE\_TEMPLATE()** [17/39]

Definition at line 989 of file barraydense-meat.hpp.

### 9.9.2.19 BDENSE\_TEMPLATE() [18/39]

Definition at line 349 of file barraydense-meat.hpp.

### **9.9.2.20** BDENSE\_TEMPLATE() [19/39]

Definition at line 341 of file barraydense-meat.hpp.

#### 9.9.2.21 BDENSE\_TEMPLATE() [20/39]

Definition at line 502 of file barraydense-meat.hpp.

#### 9.9.2.22 BDENSE\_TEMPLATE() [21/39]

Definition at line 548 of file barraydense-meat.hpp.

### 9.9.2.23 BDENSE\_TEMPLATE() [22/39]

Definition at line 552 of file barraydense-meat.hpp.

# 9.9.2.24 BDENSE\_TEMPLATE() [23/39]

Definition at line 544 of file barraydense-meat.hpp.

### 9.9.2.25 BDENSE\_TEMPLATE() [24/39]

### 9.9.2.26 BDENSE\_TEMPLATE() [25/39]

```
BDENSE_TEMPLATE (
     void ,
     clear )
```

Definition at line 896 of file barraydense-meat.hpp.

#### 9.9.2.27 BDENSE\_TEMPLATE() [26/39]

Definition at line 402 of file barraydense-meat.hpp.

#### 9.9.2.28 BDENSE\_TEMPLATE() [27/39]

# 9.9.2.29 BDENSE\_TEMPLATE() [28/39]

# 9.9.2.30 BDENSE\_TEMPLATE() [29/39]

# 9.9.2.31 BDENSE\_TEMPLATE() [30/39]

```
BDENSE_TEMPLATE (
     void ,
     reserve )
```

Definition at line 946 of file barraydense-meat.hpp.

# 9.9.2.32 BDENSE\_TEMPLATE() [31/39]

```
BDENSE_TEMPLATE (
     void ,
     resize )
```

# 9.9.2.33 BDENSE\_TEMPLATE() [32/39]

# 9.9.2.34 BDENSE\_TEMPLATE() [33/39]

# 9.9.2.35 BDENSE\_TEMPLATE() [34/39]

# 9.9.2.36 BDENSE\_TEMPLATE() [35/39]

```
BDENSE_TEMPLATE (
     void ,
     swap_rows )
```

# 9.9.2.37 BDENSE\_TEMPLATE() [36/39]

#### 9.9.2.38 BDENSE\_TEMPLATE() [37/39]

Definition at line 868 of file barraydense-meat.hpp.

# 9.9.2.39 BDENSE\_TEMPLATE() [38/39]

### 9.9.2.40 BDENSE\_TEMPLATE() [39/39]

```
BDENSE_TEMPLATE (
     void ,
     zero_row )
```

```
9.9.2.41 for()
```

```
for ( )
```

Definition at line 64 of file barraydense-meat.hpp.

el [POS(i, j)] = = BARRY\_ZERO\_DENSE)

Definition at line 663 of file barraydense-meat.hpp.

```
9.9.2.44 if() [3/4]
```

if (

```
if ( {\tt el\_colsums}~[j] ~=~ {\tt ZERO\_CELL}~)
```

# 9.9.2.45 if() [4/4]

```
if (
    el_rowsums [i] = =ZERO_CELL )
```

### 9.9.2.46 insert\_cell() [1/2]

## 9.9.2.47 insert\_cell() [2/2]

## 9.9.2.48 M()

```
bool M ( \label{eq:Array_.} \text{Array}\_. \quad \textit{M} \ )
```

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

## 9.9.2.49 printf\_barry()

```
printf_barry (
          fmt ,
          args )
```

## 9.9.2.50 resize() [1/6]

## 9.9.2.51 resize() [2/6]

## **9.9.2.52** resize() [3/6]

```
el resize ( \label{eq:N*M, ZERO_CELL} \mbox{N * $M$,}
```

## 9.9.2.53 resize() [4/6]

## 9.9.2.54 resize() [5/6]

```
el resize ( \label{eq:n_*_M_*_M_*} {\tt N_*_* M_*_*} {\tt ZERO_CELL} )
```

## 9.9.2.55 resize() [6/6]

## 9.9.2.56 rm\_cell() [1/3]

## 9.9.2.57 rm\_cell() [2/3]

## 9.9.2.58 rm\_cell() [3/3]

## 9.9.2.59 va\_end()

```
va_end (
          args )
```

#### 9.9.2.60 va\_start()

#### 9.9.3 Variable Documentation

#### 9.9.3.1 add

```
size_t const std::vector< size_t > const std::vector< size_t > bool add

Initial value:
{
    if (source.size() != target.size())
        throw std::length_error("-source- and -target- don't match on length.")
```

Definition at line 47 of file barraydense-meat.hpp.

#### 9.9.3.2 ans

```
return ans
```

Definition at line 398 of file barraydense-meat.hpp.

### 9.9.3.3 check\_bounds

```
bool check_bounds

Initial value:
{
    if (check_bounds)
    {
       out_of_range(i0,0u);
       out_of_range(i1,0u);
    }

for (size_t j = 0u; j < M; ++j)
       std::swap(el[POS(i0, j)], el[POS(i1, j)])</pre>
```

Definition at line 654 of file barraydense-meat.hpp.

## 9.9.3.4 check\_exists

```
size_t bool int check_exists

Initial value:
{
    if (check_bounds)
        out_of_range(i,j)
```

Definition at line 655 of file barraydense-meat.hpp.

#### 9.9.3.5 col

col

Definition at line 843 of file barraydense-meat.hpp.

#### 9.9.3.6 const

const

#### Initial value:

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

## 9.9.3.7 copy\_data

```
bool copy_data
```

Definition at line 156 of file barraydense-meat.hpp.

## 9.9.3.8 data

```
data = data_
```

Definition at line 334 of file barraydense-meat.hpp.

## 9.9.3.9 delete\_data

```
delete_data = delete_data_
```

Definition at line 335 of file barraydense-meat.hpp.

#### 9.9.3.10 delete\_data\_

```
bool delete_data_
```

## Initial value:

Definition at line 328 of file barraydense-meat.hpp.

#### 9.9.3.11 el

```
return el == ZERO_CELL
```

Definition at line 381 of file barraydense-meat.hpp.

#### 9.9.3.12 el\_colsums

```
el_colsums[j] = (v.value - old)
```

Definition at line 675 of file barraydense-meat.hpp.

#### 9.9.3.13 el\_rowsums

```
el_rowsums[i] = (v.value - old)
```

Definition at line 674 of file barraydense-meat.hpp.

## 9.9.3.14 else

Definition at line 670 of file barraydense-meat.hpp.

## 9.9.3.15 false

false

Definition at line 767 of file barraydense-meat.hpp.

## 9.9.3.16 i1

size\_t i1

Definition at line 721 of file barraydense-meat.hpp.

#### 9.9.3.17 j

j

Definition at line 373 of file barraydense-meat.hpp.

#### 9.9.3.18 j0

```
size_t j0
```

Definition at line 720 of file barraydense-meat.hpp.

## 9.9.3.19 j1

size\_t j1

Definition at line 721 of file barraydense-meat.hpp.

#### 9.9.3.20 M

```
M = M_{\underline{}}
```

Definition at line 57 of file barraydense-meat.hpp.

## 9.9.3.21 M\_

```
size_t M_
Initial value:
{
    std::vector< Cell_Type > el_tmp(el)
```

Definition at line 43 of file barraydense-meat.hpp.

#### 9.9.3.22 N

```
N = N_
```

Definition at line 56 of file barraydense-meat.hpp.

#### 9.9.3.23 report

```
size_t size_t bool int int* report

Initial value:
{
    if (check_bounds) {
        out_of_range(i0,j0);
        out_of_range(i1,j1);
    }

if (report != nullptr)
        (*report) = EXISTS::BOTH
```

Definition at line 724 of file barraydense-meat.hpp.

#### 9.9.3.24 return

return

Definition at line 94 of file barraydense-meat.hpp.

#### 9.9.3.25 source

```
size_t const std::vector< size_t >& source
```

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

#### 9.9.3.26 target

```
\verb|size_t| const| std::vector<|size_t| > const| std::vector<|size_t| > \& target|
```

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

#### 9.9.3.27 v

```
size_t Cell_Type v
```

Definition at line 653 of file barraydense-meat.hpp.

#### 9.9.3.28 val0

```
Cell_Type val0 = el[POS(i0,j0)]
```

Definition at line 742 of file barraydense-meat.hpp.

## 9.9.3.29 val1

```
Cell_Type val1 = el[POS(i1,j1)]
```

Definition at line 743 of file barraydense-meat.hpp.

#### 9.9.3.30 value

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

# 9.10 include/barry/barraydensecell-bones.hpp File Reference

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



## Classes

class BArrayDenseCell
 Cell\_Type, Data\_Type

### **Macros**

• #define POS(a, b) (a) + (b) \* N

## 9.10.1 Macro Definition Documentation

#### 9.10.1.1 POS

```
#define POS( \label{eq:a_posterior} \textbf{a}, \\ \textbf{b} \text{ ) (a) + (b) * N}
```

Definition at line 6 of file barraydensecell-bones.hpp.

# 9.11 include/barry/barraydensecell-meat.hpp File Reference

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



#### **Macros**

• #define POS(a, b) (a) + (b) \* dat->N

#### 9.11.1 Macro Definition Documentation

#### 9.11.1.1 POS

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

# 9.12 include/barry/barraydensecol-bones.hpp File Reference

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



#### Classes

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

#### **Macros**

```
#define POS(a, b) (b)*N + (a)
#define POS_N(a, b, c) (b)*(c) + (a)
#define ZERO_CELL static_cast<Cell_Type>(0.0)
```

## 9.12.1 Macro Definition Documentation

#### 9.12.1.1 POS

```
#define POS(  a, \\ b ) (b)*N + (a)
```

Definition at line 4 of file barraydensecol-bones.hpp.

#### 9.12.1.2 POS N

Definition at line 5 of file barraydensecol-bones.hpp.

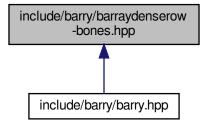
#### 9.12.1.3 ZERO\_CELL

```
#define ZERO_CELL static_cast<Cell_Type>(0.0)
```

Definition at line 6 of file barraydensecol-bones.hpp.

# 9.13 include/barry/barraydenserow-bones.hpp File Reference

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



## Classes

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

#### **Macros**

- #define POS(a, b) (b) \* N + (a)
   #define POS N(a, b, a) (b) \*(a) + (a)
- #define POS\_N(a, b, c) (b)\*(c) + (a)
- #define ZERO\_CELL static\_cast< Cell\_Type >(0.0)

## 9.13.1 Macro Definition Documentation

#### 9.13.1.1 POS

```
#define POS(  \frac{a}{b} \; ) \; \; (b) \; * \; N \; + \; (a)
```

Definition at line 4 of file barraydenserow-bones.hpp.

#### 9.13.1.2 POS N

Definition at line 5 of file barraydenserow-bones.hpp.

## 9.13.1.3 ZERO\_CELL

```
#define ZERO_CELL static_cast< Cell_Type >(0.0)
```

Definition at line 6 of file barraydenserow-bones.hpp.

## 9.14 include/barry/barrayrow-bones.hpp File Reference

#### **Classes**

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

## 9.15 include/barry/barrayrow-meat.hpp File Reference

#### **Macros**

- #define BROW\_TYPE() BArrayRow<Cell\_Type, Data\_Type>
- #define BROW\_TEMPLATE\_ARGS() < typename Cell\_Type, typename Data\_Type>
- #define BROW\_TEMPLATE(a, b) template BROW\_TEMPLATE\_ARGS() inline a BROW\_TYPE()::b

### **Functions**

- BROW\_TEMPLATE (void, operator=)(const BROW\_TYPE() &val)
- BROW\_TEMPLATE (void, operator+=)(const BROW\_TYPE() &val)
- BROW\_TEMPLATE (void, operator-=)(const BROW\_TYPE() &val)
- BROW\_TEMPLATE (void, operator\*=)(const BROW\_TYPE() &val)
- BROW\_TEMPLATE (void, operator/=)(const BROW\_TYPE() &val)

## 9.15.1 Macro Definition Documentation

## 9.15.1.1 BROW\_TEMPLATE

Definition at line 8 of file barrayrow-meat.hpp.

## 9.15.1.2 BROW\_TEMPLATE\_ARGS

```
#define BROW_TEMPLATE_ARGS() <typename Cell_Type, typename Data_Type>
```

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

## 9.15.1.3 BROW\_TYPE

```
#define BROW_TYPE( ) BArrayRow<Cell_Type, Data_Type>
```

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

### 9.15.2 Function Documentation

## 9.15.2.1 BROW\_TEMPLATE() [1/5]

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

#### 9.15.2.2 BROW\_TEMPLATE() [2/5]

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

## 9.15.2.3 BROW\_TEMPLATE() [3/5]

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

## 9.15.2.4 BROW\_TEMPLATE() [4/5]

Definition at line 55 of file barrayrow-meat.hpp.

#### 9.15.2.5 BROW\_TEMPLATE() [5/5]

```
BROW_TEMPLATE ( \mbox{void ,} \\ \mbox{operator ) const } \&
```

Definition at line 11 of file barrayrow-meat.hpp.

# 9.16 include/barry/barrayvector-bones.hpp File Reference

#### **Classes**

```
    class BArrayVector< Cell_Type, Data_Type >
        Row or column of a BArray
```

class BArrayVector\_const< Cell\_Type, Data\_Type >

## 9.17 include/barry/barrayvector-meat.hpp File Reference

## 9.18 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:

- 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\_USE\_ISFINITE When specified, it will introduce a macro that checks whether the likelihood is finite or not.
- $printf\_barry$  If not specified, will be defined as printf.
- BARRY\_DEBUG\_LEVEL, when defined, will make things verbose.
- #define BARRY\_SAFE\_EXP -100.0
- #define BARRY ISFINITE(a)
- #define BARRY\_CHECK\_SUPPORT(x, maxs)
- #define printf\_barry printf
- #define BARRY\_MAX\_NUM\_ELEMENTS static\_cast< size\_t >(std::numeric\_limits< size\_t >::max() /2u)
- template<typename Ta , typename Tb >
   using Map = std::map< Ta, Tb >

#### 9.18.1 Macro Definition Documentation

#### 9.18.1.1 BARRY CHECK SUPPORT

Definition at line 47 of file barry-configuration.hpp.

#### 9.18.1.2 BARRY\_ISFINITE

Definition at line 40 of file barry-configuration.hpp.

## 9.18.1.3 BARRY\_MAX\_NUM\_ELEMENTS

```
#define BARRY_MAX_NUM_ELEMENTS static_cast< size_t >(std::numeric_limits< size_t >::max()
/2u)
```

Definition at line 55 of file barry-configuration.hpp.

#### 9.18.1.4 BARRY\_SAFE\_EXP

```
#define BARRY_SAFE_EXP -100.0
```

Definition at line 33 of file barry-configuration.hpp.

## 9.18.1.5 printf\_barry

```
#define printf_barry printf
```

Definition at line 51 of file barry-configuration.hpp.

## 9.18.2 Typedef Documentation

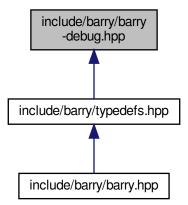
#### 9.18.2.1 Map

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

Definition at line 27 of file barry-configuration.hpp.

## 9.19 include/barry/barry-debug.hpp File Reference

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



#### **Macros**

• #define BARRY\_DEBUG\_LEVEL 0

## 9.19.1 Macro Definition Documentation

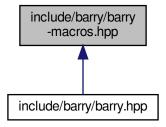
## 9.19.1.1 BARRY\_DEBUG\_LEVEL

#define BARRY\_DEBUG\_LEVEL 0

Definition at line 5 of file barry-debug.hpp.

## 9.20 include/barry/barry-macros.hpp File Reference

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



#### **Macros**

- #define BARRY\_ZERO Cell<Cell\_Type>(0.0)
- #define BARRY\_ZERO\_DENSE static\_cast<Cell\_Type>(0.0)
- #define BARRY\_ONE Cell<Cell\_Type>(1.0)
- #define BARRY\_ONE\_DENSE static\_cast<Cell\_Type>(1.0)
- #define BARRY\_UNUSED(expr) do { (void)(expr); } while (0);

#### 9.20.1 Macro Definition Documentation

### 9.20.1.1 BARRY\_ONE

```
#define BARRY_ONE CellCell_Type>(1.0)
```

Definition at line 7 of file barry-macros.hpp.

## 9.20.1.2 BARRY\_ONE\_DENSE

```
#define BARRY_ONE_DENSE static_cast<Cell_Type>(1.0)
```

Definition at line 8 of file barry-macros.hpp.

#### 9.20.1.3 BARRY\_UNUSED

Definition at line 10 of file barry-macros.hpp.

#### 9.20.1.4 BARRY\_ZERO

```
#define BARRY_ZERO Cell<Cell_Type>(0.0)
```

Definition at line 4 of file barry-macros.hpp.

## 9.20.1.5 BARRY\_ZERO\_DENSE

```
#define BARRY_ZERO_DENSE static_cast<Cell_Type>(0.0)
```

Definition at line 5 of file barry-macros.hpp.

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

```
#include <iostream>
#include <cstdarg>
#include <vector>
#include <unordered_map>
#include <functional>
#include <stdexcept>
#include <cmath>
#include <map>
#include <algorithm>
#include <utility>
#include <random>
#include <climits>
#include <cfloat>
#include <string>
#include <cstdint>
#include <memory>
#include <regex>
#include <iterator>
#include "typedefs.hpp"
#include "barry-macros.hpp"
#include "freqtable.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 "barraydense-bones.hpp"
#include "barraydensecell-bones.hpp"
#include "barraydenserow-bones.hpp"
#include "barraydensecol-bones.hpp"
#include "barraydense-meat.hpp"
#include "barraydensecell-meat.hpp"
#include "barraydense-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 dependency graph for barry.hpp:
```

## **Namespaces**

barry

barry: Your go-to motif accountant

· barry::counters

Tree class and Treelterator class.

barry::counters::network

#### **Macros**

- #define BARRY\_HPP
- #define BARRY\_VERSION\_MAYOR 0
- #define BARRY\_VERSION\_MINOR 1
- #define BARRY\_VERSION BARRY\_VERSION\_MAYOR ## . ## BARRY\_VERSION\_MINOR
- #define COUNTER FUNCTION(a)
- #define COUNTER\_LAMBDA(a)
- #define RULE\_FUNCTION(a)
- #define RULE LAMBDA(a)

#### 9.21.1 Macro Definition Documentation

## 9.21.1.1 BARRY\_HPP

```
#define BARRY_HPP
```

Definition at line 25 of file barry.hpp.

## 9.21.1.2 BARRY\_VERSION

```
#define BARRY_VERSION_BARRY_VERSION_MAYOR ## . ## BARRY_VERSION_MINOR
```

Definition at line 29 of file barry.hpp.

#### 9.21.1.3 BARRY\_VERSION\_MAYOR

```
#define BARRY_VERSION_MAYOR 0
```

Definition at line 27 of file barry.hpp.

## 9.21.1.4 BARRY\_VERSION\_MINOR

```
#define BARRY_VERSION_MINOR 1
```

Definition at line 28 of file barry.hpp.

## 9.21.1.5 COUNTER\_FUNCTION

```
#define COUNTER_FUNCTION( a )
```

#### Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
inline double (a) (const Array_Type & Array, size_t i, size_t j, Data_Type & data)\
```

Definition at line 88 of file barry.hpp.

## 9.21.1.6 COUNTER\_LAMBDA

Definition at line 91 of file barry.hpp.

#### 9.21.1.7 RULE\_FUNCTION

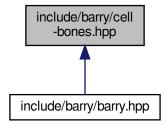
Definition at line 95 of file barry.hpp.

#### 9.21.1.8 **RULE LAMBDA**

Definition at line 98 of file barry.hpp.

# 9.22 include/barry/cell-bones.hpp File Reference

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:

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

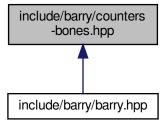
This graph shows which files directly or indirectly include this file:



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

# 9.25 include/barry/counters-bones.hpp File Reference

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

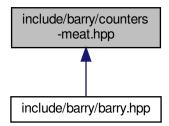


## Classes

- class Counter < Array\_Type, Data\_Type >
   A counter function based on change statistics.
- class Counters < Array\_Type, Data\_Type > Vector of counters.

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

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



#### **Macros**

- #define COUNTER\_TYPE() Counter<Array\_Type,Data\_Type>
- #define COUNTER\_TEMPLATE\_ARGS() < typename Array\_Type, typename Data\_Type >
- #define TMP\_HASHER\_CALL Hasher\_fun\_type<Array\_Type,Data\_Type>
- #define COUNTERS\_TYPE() Counters<Array\_Type,Data\_Type>
- #define COUNTERS\_TEMPLATE\_ARGS() < typename Array\_Type, typename Data\_Type>

## **Functions**

- COUNTER\_TEMPLATE (, Counter)(const Counter< Array Type
- Data\_Type init\_fun (counter\_.init\_fun)
- Data\_Type hasher\_fun (counter\_.hasher\_fun)
- Data\_Type &&counter\_\_init\_fun (std::move(counter\_\_init\_fun))
- Data\_Type &&counter\_ hasher\_fun (std::move(counter\_.hasher\_fun))
- Data Type &&counter data (std::move(counter .data))
- Data Type &&counter name (std::move(counter .name))
- Data\_Type &&counter\_ desc (std::move(counter\_.desc))

#### Move constructor.

- COUNTER\_TEMPLATE (COUNTER\_TYPE(), operator=)(const Counter< Array\_Type
- COUNTER\_TEMPLATE (COUNTER\_TYPE() &, operator=)(Counter< Array\_Type
- COUNTER TEMPLATE (double, count)(Array Type & Array

#### < Move assignment

- return count\_fun (Array, i, j, data)
- COUNTER\_TEMPLATE (double, init)(Array\_Type &Array
- return init fun (Array, i, j, data)
- COUNTER\_TEMPLATE (std::string, get\_name)() const
- COUNTER\_TEMPLATE (std::string, get\_description)() const
- COUNTER\_TEMPLATE (void, set\_hasher)(Hasher\_fun\_type< Array\_Type</li>

- COUNTER\_TEMPLATE (TMP\_HASHER\_CALL, get\_hasher)()
- COUNTERS\_TEMPLATE (, Counters)()
- COUNTERS\_TEMPLATE (COUNTER\_TYPE() &, operator[])(size\_t idx)
- Data\_Type hasher (counter\_.hasher)
- Data Type &&counters hasher (std::move(counters .hasher))
- COUNTERS TEMPLATE (COUNTERS TYPE(), operator=)(const Counters < Array Type</li>
- COUNTERS\_TEMPLATE (COUNTERS\_TYPE() &, operator=)(Counters< Array\_Type
- COUNTERS TEMPLATE (void, add counter)(Counter< Array Type</li>
- COUNTERS\_TEMPLATE (std::vector< std::string >, get\_names)() const
- COUNTERS\_TEMPLATE (std::vector< std::string >, get\_descriptions)() const
- COUNTERS TEMPLATE (std::vector< double >, gen hash)(const Array Type & array
- for (auto &c:data)
- if (add\_dims)
- if (hasher)
- if (res.size()==0u) res.push\_back(0.0)
- COUNTERS\_TEMPLATE (void, add\_hash)(Hasher\_fun\_type< Array\_Type</li>

#### **Variables**

- Data Type & counter
- Data\_Type &&counter\_ noexcept
- size\_t i = locator->second
- size\_t size\_t j
- Data\_Type fun
- Data Type counter
- · return
- Data\_Type count\_fun\_
- Data\_Type Counter\_fun\_type< Array\_Type, Data\_Type > init\_fun\_
- Data\_Type Counter\_fun\_type
   Array\_Type, Data\_Type > Hasher\_fun\_type
   Array\_Type, Data\_Type > hasher\_fun\_
- Data\_Type Counter\_fun\_type
   Array\_Type, Data\_Type > Hasher\_fun\_type
   Array\_Type, Data\_Type > Data\_Type data\_
- Data\_Type Counter\_fun\_type
   Array\_Type, Data\_Type > Hasher\_fun\_type
   Array\_Type, Data\_Type > Data\_Type std::string name
- Data\_Type Counter\_fun\_type
   Array\_Type, Data\_Type > Hasher\_fun\_type
   Array\_Type, Data\_Type > Data\_Type std::string std::string desc\_
- · bool add dims
- · return res
- Data\_Type fun\_

#### 9.26.1 Macro Definition Documentation

#### 9.26.1.1 COUNTER\_TEMPLATE

Definition at line 8 of file counters-meat.hpp.

#### 9.26.1.2 COUNTER\_TEMPLATE\_ARGS

```
#define COUNTER_TEMPLATE_ARGS() <typename Array_Type, typename Data_Type>
```

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

#### 9.26.1.3 COUNTER\_TYPE

```
#define COUNTER_TYPE( ) Counter<Array_Type,Data_Type>
```

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

#### 9.26.1.4 COUNTERS\_TEMPLATE

Definition at line 129 of file counters-meat.hpp.

### 9.26.1.5 COUNTERS\_TEMPLATE\_ARGS

```
#define COUNTERS_TEMPLATE_ARGS() <typename Array_Type, typename Data_Type>
```

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

#### 9.26.1.6 COUNTERS\_TYPE

```
#define COUNTERS_TYPE( ) Counters<Array_Type,Data_Type>
```

Definition at line 125 of file counters-meat.hpp.

## 9.26.1.7 TMP\_HASHER\_CALL

```
#define TMP_HASHER_CALL Hasher_fun_type<Array_Type,Data_Type>
```

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

#### 9.26.2 Function Documentation

#### 9.26.2.1 count fun()

## 9.26.2.2 COUNTER\_TEMPLATE() [1/9]

```
COUNTER_TEMPLATE (

Counter ) const
```

## 9.26.2.3 COUNTER\_TEMPLATE() [2/9]

## 9.26.2.4 COUNTER\_TEMPLATE() [3/9]

## 9.26.2.5 COUNTER\_TEMPLATE() [4/9]

 $< {\sf Move \ assignment}$ 

## **9.26.2.6 COUNTER\_TEMPLATE()** [5/9]

## 9.26.2.7 **COUNTER\_TEMPLATE()** [6/9]

```
COUNTER_TEMPLATE (
          std::string ,
          get_description ) const
```

Definition at line 107 of file counters-meat.hpp.

## 9.26.2.8 **COUNTER\_TEMPLATE()** [7/9]

```
COUNTER_TEMPLATE (
          std::string ,
          get_name ) const
```

Definition at line 103 of file counters-meat.hpp.

### 9.26.2.9 COUNTER\_TEMPLATE() [8/9]

```
COUNTER_TEMPLATE (

TMP_HASHER_CALL ,

qet_hasher )
```

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

#### 9.26.2.10 COUNTER\_TEMPLATE() [9/9]

## 9.26.2.11 COUNTERS\_TEMPLATE() [1/9]

```
COUNTERS_TEMPLATE (

Counters )
```

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

#### 9.26.2.12 COUNTERS\_TEMPLATE() [2/9]

```
COUNTERS_TEMPLATE (

COUNTER_TYPE() & ,

operator [])
```

Definition at line 134 of file counters-meat.hpp.

## 9.26.2.13 COUNTERS\_TEMPLATE() [3/9]

```
COUNTERS_TEMPLATE (

COUNTERS_TYPE() & ,

operator )
```

## 9.26.2.14 COUNTERS\_TEMPLATE() [4/9]

```
COUNTERS_TEMPLATE (

COUNTERS_TYPE() ,

operator ) const
```

## 9.26.2.15 COUNTERS\_TEMPLATE() [5/9]

```
COUNTERS_TEMPLATE (
          std::vector< double > ,
          gen_hash ) const &
```

#### 9.26.2.16 COUNTERS\_TEMPLATE() [6/9]

```
COUNTERS_TEMPLATE (
          std::vector< std::string > ,
          get_descriptions ) const
```

Definition at line 212 of file counters-meat.hpp.

## 9.26.2.17 COUNTERS\_TEMPLATE() [7/9]

```
COUNTERS_TEMPLATE (
          std::vector< std::string > ,
          get_names ) const
```

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

## 9.26.2.18 COUNTERS\_TEMPLATE() [8/9]

## 9.26.2.19 **COUNTERS\_TEMPLATE()** [9/9]

## 9.26.2.20 data()

### 9.26.2.21 desc()

Move constructor.

Definition at line 32 of file counters-meat.hpp.

### 9.26.2.22 for()

```
for ( auto &c:data )
```

Definition at line 231 of file counters-meat.hpp.

## 9.26.2.23 hasher() [1/2]

Definition at line 141 of file counters-meat.hpp.

## 9.26.2.24 hasher() [2/2]

Definition at line 144 of file counters-meat.hpp.

## 9.26.2.25 hasher\_fun() [1/2]

Definition at line 13 of file counters-meat.hpp.

#### 9.26.2.26 hasher\_fun() [2/2]

## 9.26.2.27 if() [1/3]

```
if (
    add_dims )
```

Definition at line 246 of file counters-meat.hpp.

## 9.26.2.28 if() [2/3]

```
if ( hasher )
```

Definition at line 253 of file counters-meat.hpp.

```
9.26.2.29 if() [3/3]
```

```
if ( {\tt res.} \quad {\tt size() = =0u \ )}
```

### 9.26.2.30 init\_fun() [1/3]

## 9.26.2.31 init\_fun() [2/3]

## **9.26.2.32** init\_fun() [3/3]

### 9.26.2.33 name()

## 9.26.3 Variable Documentation

#### 9.26.3.1 add\_dims

```
bool add_dims
```

## Initial value:

```
std::vector<double> res
```

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

## 9.26.3.2 count\_fun\_

```
Data_Type count_fun_
```

Definition at line 179 of file counters-meat.hpp.

#### 9.26.3.3 counter

```
Data_Type counter

Initial value:
{
```

data.push\_back(counter)

Definition at line 170 of file counters-meat.hpp.

## 9.26.3.4 counter\_

```
Data_Type & counter_

Initial value:
{
    if (this != &counter_) {
        this->count_fun = counter_.count_fun;
        this->init_fun = counter_.init_fun;
        this->hasher_fun = counter_.hasher_fun;

        this->data = counter_.data;
        this->name = counter_.name;
        this->desc = counter_.desc;
    }
    return *this
```

Definition at line 12 of file counters-meat.hpp.

## 9.26.3.5 data\_

)

```
Data_Rule_Dyn_Type Data_Rule_Dyn_Type data_
Initial value:
{
    rules_dyn->add_rule(
        rule_fun_,
        data_
```

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

#### 9.26.3.6 desc\_

```
Data_Type Counter_fun_type<Array_Type,Data_Type> Hasher_fun_type<Array_Type,Data_Type> Data←
_Type std::string std::string desc_
```

#### Initial value:

```
data.push_back(Counter<Array_Type,Data_Type>(
    count_fun_,
    init_fun_,
    hasher_fun_,
    data_,
    name_,
    desc_
))
```

Definition at line 184 of file counters-meat.hpp.

#### 9.26.3.7 fun

```
Data_Type fun

Initial value:
{
    hasher_fun = fun
```

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

#### 9.26.3.8 fun

```
Data_Type fun_
Initial value:
```

```
{
    hasher = fun_
```

Definition at line 268 of file counters-meat.hpp.

## 9.26.3.9 hasher\_fun\_

Data\_Type Counter\_fun\_type<Array\_Type,Data\_Type> Hasher\_fun\_type<Array\_Type,Data\_Type> hasher← \_fun\_

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

#### 9.26.3.10 i

```
const std::vector< double > size_t i = locator->second
```

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

#### 9.26.3.11 init fun

```
Data_Type Counter_fun_type<Array_Type,Data_Type> init_fun_
```

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

#### 9.26.3.12 j

```
size_t size_t j

Initial value:
{
    if (count_fun == nullptr)
        return 0.0
```

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

#### 9.26.3.13 name\_

```
Data_Type Counter_fun_type<Array_Type,Data_Type> Hasher_fun_type<Array_Type,Data_Type> Data←
_Type std::string name_
```

Definition at line 183 of file counters-meat.hpp.

#### 9.26.3.14 noexcept

```
Data_Type &&counters_ noexcept

Initial value:
{
    if (this != &counter_)
    {
        this->data = std::move(counter_.data);

        this->count_fun = std::move(counter_.count_fun);
        this->init_fun = std::move(counter_.init_fun);
        this->hasher_fun = std::move(counter_.hasher_fun);

        this->name = std::move(counter_.name);
        this->desc = std::move(counter_.desc);
    }
}
```

Definition at line 26 of file counters-meat.hpp.

#### 9.26.3.15 res

return res

Definition at line 263 of file counters-meat.hpp.

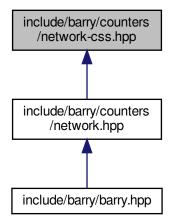
#### 9.26.3.16 return

return

Definition at line 175 of file counters-meat.hpp.

# 9.27 include/barry/counters/network-css.hpp File Reference

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



#### **Macros**

- #define CSS\_SIZE()
- #define CSS\_CASE\_TRUTH() if ((i < n) && (j < n))</li>
- #define CSS\_TRUE\_CELLS()
- #define CSS\_CASE\_PERCEIVED() else if ((( $i \ge s$ ) && (i < e)) & (( $j \ge s$ ) && (j < e)))
- #define CSS\_PERCEIVED\_CELLS()
- #define CSS\_CASE\_ELSE()
- #define CSS\_CHECK\_SIZE\_INIT()
- #define CSS\_CHECK\_SIZE()
- #define CSS\_APPEND(name)
- #define CSS\_NET\_COUNTER\_LAMBDA\_INIT()

#### **Functions**

```
• template<typename Tnet = Network>
  void counter css partially false recip commi (NetCounters< Tnet > *counters, size t netsize, const std ←
  ::vector< size t > &end )
     Counts errors of commission.
• template<typename Tnet = Network>
  void counter css partially false recip omiss (NetCounters< Tnet > *counters, size t netsize, const std↔
  ::vector< size t > &end )
     Counts errors of omission.
• template<typename Tnet = Network>
  void counter css completely false recip comiss (NetCounters< Tnet > *counters, size t netsize, const
  std::vector < size_t > &end_)
     Counts completely false reciprocity (comission)
template<typename Tnet = Network>
  void counter css completely false recip omiss (NetCounters< Tnet > *counters, size t netsize, const
  std::vector< size_t > &end_)
     Counts completely false reciprocity (omission)
template<typename Tnet = Network>
  void counter css mixed recip (NetCounters< Tnet > *counters, size t netsize, const std::vector< size t >
  &end )
     Counts mixed reciprocity errors.
• template<typename Tnet = Network>
  void counter_css_census01 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end )
• template<typename Tnet = Network>
  void counter_css_census02 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size t >
  &end )

    template<typename Tnet = Network>

  void counter_css_census03 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end )
template<typename Tnet = Network>
  void counter_css_census04 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end_)
template<typename Tnet = Network>
  void counter_css_census05 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end )
template<typename Tnet = Network>
  void counter css census06 (NetCounters< Tnet > *counters, size t netsize, const std::vector< size t >
  &end )
template<typename Tnet = Network>
  void counter_css_census07 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end_)
template<typename Tnet = Network>
  void counter_css_census08 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end )
template<typename Tnet = Network>
  void counter css census09 (NetCounters< Tnet > *counters, size t netsize, const std::vector< size t >
  &end )
template<typename Tnet = Network>
  void counter css census10 (NetCounters< Tnet > *counters, size t netsize, const std::vector< size t >
  &end )
```

#### 9.27.1 Macro Definition Documentation

#### 9.27.1.1 CSS\_APPEND

Definition at line 42 of file network-css.hpp.

### 9.27.1.2 CSS\_CASE\_ELSE

```
#define CSS_CASE_ELSE( )
```

Definition at line 27 of file network-css.hpp.

#### 9.27.1.3 CSS\_CASE\_PERCEIVED

Definition at line 20 of file network-css.hpp.

### 9.27.1.4 CSS\_CASE\_TRUTH

```
#define CSS_CASE_TRUTH( ) if ((i < n) && (j < n))
```

Definition at line 13 of file network-css.hpp.

#### 9.27.1.5 CSS\_CHECK\_SIZE

```
#define CSS_CHECK_SIZE( )
```

#### Value:

```
for (size_t i = 0u; i < end_.size(); ++i) {\
  if (i == 0u) continue; \
  else if (end_[i] < end_[i-1u]) \
     throw std::logic_error("Endpoints should be specified in order.");}</pre>
```

Definition at line 37 of file network-css.hpp.

### 9.27.1.6 CSS\_CHECK\_SIZE\_INIT

Definition at line 31 of file network-css.hpp.

#### 9.27.1.7 CSS\_NET\_COUNTER\_LAMBDA\_INIT

Definition at line 49 of file network-css.hpp.

### 9.27.1.8 CSS\_PERCEIVED\_CELLS

```
#define CSS_PERCEIVED_CELLS( )

Value:
    double tji = static_cast<double>(Array(j - s, i - s, false)); \
    double pji = static_cast<double>(Array(j, i, false)); \
    double tij = static_cast<double>(Array(i - s, j - s, false));
```

Definition at line 21 of file network-css.hpp.

#### 9.27.1.9 CSS\_SIZE

```
#define CSS_SIZE( )

Value:
    size_t n = data.indices[0u]; \
    size_t s = data.indices[1u]; \
    size_t e = data.indices[2u];
```

Definition at line 7 of file network-css.hpp.

### 9.27.1.10 CSS\_TRUE\_CELLS

```
#define CSS_TRUE_CELLS( )

Value:
    double tji = static_cast<double>(Array(j, i, false)); \
    double pij = static_cast<double>(Array(i + s, j + s, false)); \
    double pji = static_cast<double>(Array(j + s, i + s, false));
```

Definition at line 14 of file network-css.hpp.

#### 9.27.2 Function Documentation

#### 9.27.2.1 counter\_css\_census01()

Definition at line 275 of file network-css.hpp.

#### 9.27.2.2 counter\_css\_census02()

Definition at line 325 of file network-css.hpp.

#### 9.27.2.3 counter\_css\_census03()

Definition at line 364 of file network-css.hpp.

#### 9.27.2.4 counter\_css\_census04()

Definition at line 403 of file network-css.hpp.

#### 9.27.2.5 counter\_css\_census05()

Definition at line 442 of file network-css.hpp.

#### 9.27.2.6 counter\_css\_census06()

Definition at line 481 of file network-css.hpp.

### 9.27.2.7 counter\_css\_census07()

Definition at line 520 of file network-css.hpp.

#### 9.27.2.8 counter\_css\_census08()

Definition at line 559 of file network-css.hpp.

#### 9.27.2.9 counter\_css\_census09()

Definition at line 598 of file network-css.hpp.

#### 9.27.2.10 counter\_css\_census10()

Definition at line 637 of file network-css.hpp.

### 9.27.2.11 counter\_css\_completely\_false\_recip\_comiss()

Counts completely false reciprocity (comission)

Definition at line 154 of file network-css.hpp.

#### 9.27.2.12 counter\_css\_completely\_false\_recip\_omiss()

Counts completely false reciprocity (omission)

Definition at line 194 of file network-css.hpp.

#### 9.27.2.13 counter\_css\_mixed\_recip()

Counts mixed reciprocity errors.

Definition at line 234 of file network-css.hpp.

#### 9.27.2.14 counter\_css\_partially\_false\_recip\_commi()

Counts errors of commission.

#### **Parameters**

netsize	Size of the reference (true) network
end←	Vector indicating one past the ending index of each network. (see details)
_	

The  ${\tt end\_parameter}$  should be of length N of  ${\tt networks-1}$ . It is assumed that the first network ends at  ${\tt netsize}$ .

Definition at line 63 of file network-css.hpp.

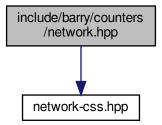
### 9.27.2.15 counter\_css\_partially\_false\_recip\_omiss()

Counts errors of omission.

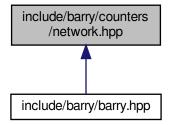
Definition at line 110 of file network-css.hpp.

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

```
#include "network-css.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 size\_t 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)
- #define NETWORKDENSE COUNTER LAMBDA(a)

#### Macros for defining rules

- #define NETWORK RULE(a)
- #define NETWORK RULE LAMBDA(a)

#### **Functions**

```
    template<typename Tnet = Network>
        void counter_edges (NetCounters< Tnet > *counters)
```

Number of edges.

template<typename Tnet = Network>
 void counter\_isolates (NetCounters< Tnet > \*counters)

Number of isolated vertices.

- template<> void counter\_isolates (NetCounters< NetworkDense > \*counters)
- template<typename Tnet = Network>
   void counter\_mutual (NetCounters< Tnet > \*counters)

Number of mutual ties.

- template<typename Tnet = Network>
  - void counter\_istar2 (NetCounters< Tnet > \*counters)
- template<> void counter\_istar2 (NetCounters< NetworkDense > \*counters)
- template<typename Tnet = Network>
  - $void\ counter\_ostar2\ (NetCounters < Tnet > *counters)$
- template<> void counter\_ostar2 (NetCounters< NetworkDense > \*counters)
- template<typename Tnet = Network>
  - void counter\_ttriads (NetCounters < Tnet > \*counters)
- template<> void counter\_ttriads (NetCounters< NetworkDense > \*counters)
- template<typename Tnet = Network>
  - void counter ctriads (NetCounters< Tnet > \*counters)
- template<> void counter\_ctriads (NetCounters< NetworkDense > \*counters)
- template < typename Tnet = Network >
   void counter\_density (NetCounters < Tnet > \*counters)

```
• template<typename Tnet = Network>
  void counter_idegree15 (NetCounters< Tnet > *counters)

    template<> void counter_idegree15 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter odegree15 (NetCounters< Tnet > *counters)

    template<> void counter_odegree15 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter_absdiff (NetCounters< Tnet > *counters, size_t attr_id, double alpha=1.0)
     Sum of absolute attribute difference between ego and alter.
• template<typename Tnet = Network>
  void counter diff (NetCounters < Tnet > *counters, size t 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)

• template<typename Tnet = Network>
  void counter_nodeicov (NetCounters< Tnet > *counters, size_t attr_id)

    template<tvpename Tnet = Network>

  void counter_nodeocov (NetCounters< Tnet > *counters, size_t attr_id)

    template<typename Tnet = Network>

  void counter_nodecov (NetCounters< Tnet > *counters, size_t attr_id)
template<typename Tnet = Network>
  void counter_nodematch (NetCounters< Tnet > *counters, size_t attr_id)
• template<typename Tnet = Network>
  void counter idegree (NetCounters< Tnet > *counters, std::vector< size t > d)
     Counts number of vertices with a given in-degree.

    template<> void counter_idegree (NetCounters< NetworkDense > *counters, std::vector< size_t > d)

• template<typename Tnet = Network>
  void counter_odegree (NetCounters< Tnet > *counters, std::vector< size_t > d)
     Counts number of vertices with a given out-degree.

    template<> void counter odegree (NetCounters< NetworkDense > *counters, std::vector< size t > d)

template<typename Tnet = Network>
  void\ counter\_degree\ (NetCounters < Tnet > *counters,\ std::vector < size\_t > 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 > ).
```

template < typename Tnet = Network >
 void rules\_zerodiag (NetRules < Tnet > \*rules)
 Number of edges.

#### Convenient typedefs for network objects.

- #define BARRY\_ZERO\_NETWORK 0.0
- #define BARRY\_ZERO\_NETWORK\_DENSE 0
- typedef BArray< double, NetworkData > Network
- typedef BArrayDense< int, NetworkData > NetworkDense
- template<typename Tnet = Network>
   using NetCounter = Counter< Tnet, NetCounterData >
- template < typename Tnet = Network >
   using NetCounters = Counters < Tnet, NetCounterData >

```
    template < typename Tnet = Network>
        using NetSupport = Support < Tnet, NetCounterData >
    template < typename Tnet = Network>
        using NetStatsCounter = StatsCounter < Tnet, NetCounterData >
    template < typename Tnet >
        using NetModel = Model < Tnet, NetCounterData >
    template < typename Tnet = Network>
        using NetRule = Rule < Tnet, bool >
    template < typename Tnet = Network>
        using NetRules = Rules < Tnet, bool >
```

#### 9.28.1 Macro Definition Documentation

### 9.28.1.1 BARRY\_ZERO\_NETWORK

```
#define BARRY_ZERO_NETWORK 0.0
```

Definition at line 85 of file network.hpp.

### 9.28.1.2 BARRY\_ZERO\_NETWORK\_DENSE

```
#define BARRY_ZERO_NETWORK_DENSE 0
```

Definition at line 86 of file network.hpp.

#### 9.28.1.3 NET\_C\_DATA\_IDX

Definition at line 74 of file network.hpp.

### 9.28.1.4 NET\_C\_DATA\_NUM

Definition at line 75 of file network.hpp.

### 9.28.1.5 NETWORK\_COUNTER

Function for definition of a network counter function

Definition at line 114 of file network.hpp.

#### 9.28.1.6 NETWORK COUNTER LAMBDA

Lambda function for definition of a network counter function

Definition at line 119 of file network.hpp.

#### 9.28.1.7 **NETWORK RULE**

#define NETWORK\_RULE(

```
Value:
template<typename Tnet = Network>\
inline bool (a) (const Tnet & Array, size_t i, size_t j, bool & data)
```

Function for definition of a network counter function

Definition at line 133 of file network.hpp.

### 9.28.1.8 NETWORK\_RULE\_LAMBDA

Lambda function for definition of a network counter function

Definition at line 138 of file network.hpp.

### 9.28.1.9 NETWORKDENSE\_COUNTER\_LAMBDA

Definition at line 123 of file network.hpp.

### 9.28.2 Typedef Documentation

#### 9.28.2.1 NetCounter

```
template<typename Tnet = Network>
using NetCounter = Counter<Tnet, NetCounterData >
```

Definition at line 89 of file network.hpp.

### 9.28.2.2 NetCounters

```
template<typename Tnet = Network>
using NetCounters = Counters<Tnet, NetCounterData>
```

Definition at line 92 of file network.hpp.

#### 9.28.2.3 NetModel

```
template<typename Tnet >
using NetModel = Model<Tnet, NetCounterData>
```

Definition at line 101 of file network.hpp.

#### 9.28.2.4 NetRule

```
template<typename Tnet = Network>
using NetRule = Rule<Tnet, bool>
```

Definition at line 104 of file network.hpp.

#### 9.28.2.5 NetRules

```
template<typename Tnet = Network>
using NetRules = Rules<Tnet, bool>
```

Definition at line 107 of file network.hpp.

#### 9.28.2.6 NetStatsCounter

```
template<typename Tnet = Network>
using NetStatsCounter = StatsCounter<Tnet, NetCounterData>
```

Definition at line 98 of file network.hpp.

### 9.28.2.7 NetSupport

```
template<typename Tnet = Network>
using NetSupport = Support<Tnet, NetCounterData >
```

Definition at line 95 of file network.hpp.

#### 9.28.2.8 Network

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

Definition at line 82 of file network.hpp.

#### 9.28.2.9 NetworkDense

```
typedef BArrayDense<int, NetworkData> NetworkDense
```

Definition at line 83 of file network.hpp.

#### 9.28.3 Function Documentation

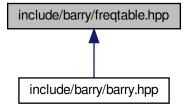
#### 9.28.3.1 rules\_zerodiag()

Number of edges.

Definition at line 1381 of file network.hpp.

# 9.29 include/barry/freqtable.hpp File Reference

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

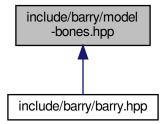


### Classes

class FreqTable < T >
 Frequency table of vectors.

# 9.30 include/barry/model-bones.hpp File Reference

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



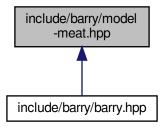
#### Classes

class Model < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type >

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

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

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



#### **Macros**

- #define MODEL TYPE()
- #define MODEL TEMPLATE ARGS()
- #define MODEL\_TEMPLATE(a, b) template MODEL\_TEMPLATE\_ARGS() inline a MODEL\_TYPE()::b

#### **Functions**

- double update\_normalizing\_constant (const double \*params, const double \*support, size\_t k, size\_t n)
- double likelihood\_ (const double \*stats\_target, const std::vector< double > &params, const double normalizing constant, size t n params, bool log =false)
- MODEL\_TEMPLATE (void, store\_psets)() noexcept
- MODEL\_TEMPLATE (std::vector< double >, gen\_key)(const Array\_Type &Array\_)
- MODEL\_TEMPLATE (void, add\_counter)(Counter< Array\_Type
- MODEL\_TEMPLATE (void, set\_counters)(Counters < Array\_Type</li>
- support\_fun set\_counters (counters)
- MODEL TEMPLATE (void, add hasher)(Hasher fun type< Array Type
- MODEL TEMPLATE (void, add rule)(Rule< Array Type
- MODEL\_TEMPLATE (void, set\_rules)(Rules< Array\_Type
- support fun set rules (rules)
- MODEL\_TEMPLATE (void, add\_rule\_dyn)(Rule < Array\_Type</li>
- MODEL\_TEMPLATE (void, set\_rules\_dyn)(Rules < Array\_Type</li>
- support\_fun set\_rules\_dyn (rules\_dyn)
- MODEL\_TEMPLATE (size\_t, add\_array)(const Array\_Type &Array\_
- if (transform\_model\_fun) = transform\_model\_fun(&tmp\_counts[0u], tmp\_counts.size())
- else stats\_target push\_back (counter\_fun.count\_all())

```
    if (force new|(locator==keys2support.end()))
```

- arrays2support push back (locator->second)
- return arrays2support size () 1u
- MODEL TEMPLATE (double, likelihood)(const std
- MODEL TEMPLATE (double, likelihood total)(const std
- MODEL TEMPLATE (double, get norm const)(const std
- MODEL\_TEMPLATE (const std::vector< Array\_Type > \*, get\_pset)(const size\_t &i)
- MODEL TEMPLATE (const std::vector< double > \*, get pset stats)(const size t &i)
- MODEL\_TEMPLATE (void, print\_stats)(size\_t i) const
- MODEL\_TEMPLATE (size\_t, size)() const noexcept
- MODEL TEMPLATE (size t, size unique)() const noexcept
- MODEL\_TEMPLATE (size\_t, nterms)() const noexcept
- MODEL TEMPLATE (size t, nrules)() const noexcept
- MODEL\_TEMPLATE (size\_t, nrules\_dyn)() const noexcept
- MODEL TEMPLATE (size t, support size)() const noexcept
- MODEL\_TEMPLATE (std::vector< std::string >, colnames)() const
- MODEL TEMPLATE (Array Type, sample)(const Array Type & Array
- if (locator==keys2support.end())
- std::uniform\_real\_distribution urand (0, 1)
- if ((probs.size() > 0u) &&(vec equal approx(params, params last[a])))
- std::vector< double > temp\_stats (params.size())
- for (size t array=0u;array< probs.size();++array)</li>
- MODEL TEMPLATE (double, conditional prob)(const Array Type & Array
- A insert cell (i, j, A.default val(), true, false)
- std::vector< double > tmp\_counts (counters->size())
- return (1.0+std::exp(-vec inner prod < double >(&params[0u], &tmp counts[0u], params.size())))
- MODEL\_TEMPLATE (const std::mt19937 \*, get\_rengine)() const
- MODEL TEMPLATE (std::vector< std::vector< double > > \*, get stats target)()
- MODEL TEMPLATE (std::vector< std::vector< double >> \*, get stats support)()
- MODEL\_TEMPLATE (std::vector< size\_t > \*, get\_arrays2support)()
- MODEL TEMPLATE (std::vector < std::vector < Array Type > > \*, get pset arrays)()
- MODEL\_TEMPLATE (std::vector< std::vector< double >> \*, get\_pset\_stats)()
- MODEL TEMPLATE (std::vector< std::vector< double >> \*, get\_pset\_probs)()
- MODEL TEMPLATE (void, set transform model)(std

#### **Variables**

- Data\_Counter\_Type & counter
- return
- Data Counter Type count fun
- Data\_Counter\_Type Counter\_fun\_type< Array\_Type, Data\_Counter\_Type > init\_fun\_
- Data\_Counter\_Type Counter\_fun\_type < Array\_Type, Data\_Counter\_Type > Data\_Counter\_Type data\_
- Data Counter Type \* counters
- Data\_Counter\_Type fun\_
- · Data Rule Type & rules
- Data Rule Type \* rules
- this delete rules = false
- Data Rule Dyn Type rule fun
- this rules\_dyn = rules\_
- this delete\_rules\_dyn = false
- bool force new
- std::vector< double > key = counters->gen hash(Array )
- MapVec\_type< double, size\_t >::const\_iterator locator = keys2support.find(key)
- stats\_support\_n\_arrays [locator->second]

```
const std::vector< double > & params
size_t i = locator->second
size_t a = arrays2support[i]
double r = urand(*rengine)
double cumprob = 0.0
size_t k = params.size()
size_t j = 0u
std::vector< double > & probs = pset_probs[a]
else
const std::vector< double > & stats = pset_stats[a]
int i_matches = -1
return this pset_arrays [a][j]
template Data_Counter_Type
template Data_Rule_Type
```

#### 9.31.1 Macro Definition Documentation

#### 9.31.1.1 MODEL\_TEMPLATE

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

#### 9.31.1.2 MODEL\_TEMPLATE\_ARGS

Definition at line 120 of file model-meat.hpp.

#### 9.31.1.3 MODEL\_TYPE

```
template Data_Rule_Dyn_Type * MODEL_TYPE( )

Value:
         Model<Array_Type, Data_Counter_Type, Data_Rule_Type, \
         Data_Rule_Dyn_Type>
```

Definition at line 117 of file model-meat.hpp.

#### 9.31.2 Function Documentation

```
9.31.2.1 for()
```

```
for ( )
```

Definition at line 1307 of file model-meat.hpp.

```
9.31.2.2 if() [1/4]
```

```
if (
     (probs.size() > Ou) &&(vec_equal_approx(params, params_last[a])) )
```

Definition at line 1290 of file model-meat.hpp.

```
9.31.2.3 if() [2/4]
```

```
if (
          force_new| locator==keys2support.end() )
```

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

Definition at line 449 of file model-meat.hpp.

```
9.31.2.4 if() [3/4]
```

```
if (
    locator = = keys2support.end() )
```

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

Definition at line 1192 of file model-meat.hpp.

```
9.31.2.5 if() [4/4]
```

```
if (
    transform_model_fun ) = transform_model_fun(&tmp_counts[Ou], tmp_counts.size())
```

Definition at line 434 of file model-meat.hpp.

### 9.31.2.6 insert\_cell()

#### 9.31.2.7 likelihood\_()

Definition at line 59 of file model-meat.hpp.

#### 9.31.2.8 MODEL\_TEMPLATE() [1/33]

```
MODEL_TEMPLATE ( \label{eq:array_Type} \mbox{ Array_Type ,} \\ \mbox{sample ) const &} \mbox{}
```

### 9.31.2.9 MODEL\_TEMPLATE() [2/33]

Definition at line 1376 of file model-meat.hpp.

### 9.31.2.10 MODEL\_TEMPLATE() [3/33]

Definition at line 918 of file model-meat.hpp.

### 9.31.2.11 MODEL\_TEMPLATE() [4/33]

Definition at line 930 of file model-meat.hpp.

### 9.31.2.12 MODEL\_TEMPLATE() [5/33]

### 9.31.2.13 MODEL\_TEMPLATE() [6/33]

Definition at line 882 of file model-meat.hpp.

# **9.31.2.14 MODEL\_TEMPLATE()** [7/33]

Definition at line 561 of file model-meat.hpp.

#### 9.31.2.15 MODEL\_TEMPLATE() [8/33]

Definition at line 816 of file model-meat.hpp.

### 9.31.2.16 MODEL\_TEMPLATE() [9/33]

#### 9.31.2.17 MODEL\_TEMPLATE() [10/33]

Definition at line 1056 of file model-meat.hpp.

### 9.31.2.18 MODEL\_TEMPLATE() [11/33]

Definition at line 1063 of file model-meat.hpp.

#### 9.31.2.19 MODEL\_TEMPLATE() [12/33]

Definition at line 1046 of file model-meat.hpp.

#### 9.31.2.20 MODEL\_TEMPLATE() [13/33]

Definition at line 1031 of file model-meat.hpp.

### 9.31.2.21 MODEL\_TEMPLATE() [14/33]

Definition at line 1038 of file model-meat.hpp.

### 9.31.2.22 MODEL\_TEMPLATE() [15/33]

Definition at line 1070 of file model-meat.hpp.

#### 9.31.2.23 MODEL\_TEMPLATE() [16/33]

Definition at line 304 of file model-meat.hpp.

#### 9.31.2.24 MODEL\_TEMPLATE() [17/33]

Definition at line 1411 of file model-meat.hpp.

### **9.31.2.25** MODEL\_TEMPLATE() [18/33]

```
MODEL_TEMPLATE (
          std::vector< std::string > ,
          colnames ) const
```

Definition at line 1082 of file model-meat.hpp.

#### **9.31.2.26** MODEL\_TEMPLATE() [19/33]

Definition at line 1416 of file model-meat.hpp.

### 9.31.2.27 MODEL\_TEMPLATE() [20/33]

```
MODEL_TEMPLATE (
          std::vector< std::vector< double > > * ,
          get_pset_probs )
```

Definition at line 1424 of file model-meat.hpp.

#### 9.31.2.28 MODEL\_TEMPLATE() [21/33]

```
MODEL_TEMPLATE (
          std::vector< std::vector< double > > * ,
          get_pset_stats )
```

Definition at line 1420 of file model-meat.hpp.

# 9.31.2.29 MODEL\_TEMPLATE() [22/33]

```
MODEL_TEMPLATE (
          std::vector< std::vector< double > > * ,
          get_stats_support )
```

Definition at line 1406 of file model-meat.hpp.

### 9.31.2.30 MODEL\_TEMPLATE() [23/33]

```
MODEL_TEMPLATE (
          std::vector< std::vector< double > > * ,
          get_stats_target )
```

Definition at line 1401 of file model-meat.hpp.

### 9.31.2.31 MODEL\_TEMPLATE() [24/33]

### 9.31.2.32 MODEL\_TEMPLATE() [25/33]

```
MODEL_TEMPLATE (
     void ,
     add_hasher )
```

## 9.31.2.33 MODEL\_TEMPLATE() [26/33]

#### 9.31.2.34 MODEL\_TEMPLATE() [27/33]

#### **9.31.2.35 MODEL\_TEMPLATE()** [28/33]

Definition at line 941 of file model-meat.hpp.

### 9.31.2.36 MODEL\_TEMPLATE() [29/33]

### **9.31.2.37** MODEL\_TEMPLATE() [30/33]

#### 9.31.2.38 MODEL\_TEMPLATE() [31/33]

#### 9.31.2.39 MODEL\_TEMPLATE() [32/33]

Definition at line 1428 of file model-meat.hpp.

### 9.31.2.40 MODEL\_TEMPLATE() [33/33]

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

### 9.31.2.41 push\_back() [1/2]

### 9.31.2.42 push\_back() [2/2]

### 9.31.2.43 return()

```
return (  1.0+ \ std::exp-vec\_inner\_prod < \ double > (\&params[0u], \ \&tmp\_counts[0u], \ params. \leftarrow \\ size()) \ )
```

### 9.31.2.44 set\_counters()

### 9.31.2.45 set\_rules()

#### 9.31.2.46 set\_rules\_dyn()

#### 9.31.2.47 size()

```
return arrays2support size ( )
```

### 9.31.2.48 temp\_stats()

```
std::vector< double > temp_stats (
    params. size() )
```

### 9.31.2.49 tmp\_counts()

### 9.31.2.50 update\_normalizing\_constant()

Definition at line 9 of file model-meat.hpp.

#### 9.31.2.51 urand()

```
std::uniform_real_distribution urand ( \begin{array}{c} 0 \text{ ,} \\ 1 \end{array} )
```

#### 9.31.3 Variable Documentation

#### 9.31.3.1 a

```
size_t a = arrays2support[i]
```

Definition at line 1278 of file model-meat.hpp.

### 9.31.3.2 count\_fun\_

```
Data_Counter_Type count_fun_
```

Definition at line 319 of file model-meat.hpp.

#### 9.31.3.3 counter

```
Data_Counter_Type& counter

Initial value:
{
    counters->add_counter(counter, Data_Counter_Type())
```

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

#### 9.31.3.4 counters\_

```
Data_Counter_Type* counters_
Initial value:
{
    if (delete_counters) {
        delete counters;
        delete_counters = false;
    }
    this->counters = counters_
```

Definition at line 335 of file model-meat.hpp.

#### 9.31.3.5 cumprob

```
double cumprob = 0.0
```

Definition at line 1283 of file model-meat.hpp.

#### 9.31.3.6 data\_

```
Data_Rule_Dyn_Type Data_Rule_Dyn_Type data_
Initial value:
{
    counters->add_counter(
        count_fun_,
        init_fun_,
        data_
```

Definition at line 321 of file model-meat.hpp.

### 9.31.3.7 Data\_Counter\_Type

```
template Data_Counter_Type
```

Definition at line 1396 of file model-meat.hpp.

### 9.31.3.8 Data\_Rule\_Type

```
template Data_Rule_Type
```

Definition at line 1396 of file model-meat.hpp.

### 9.31.3.9 delete\_rules

```
this delete_rules = false
```

Definition at line 378 of file model-meat.hpp.

# 9.31.3.10 delete\_rules\_dyn

```
this delete_rules_dyn = false
```

Definition at line 417 of file model-meat.hpp.

#### 9.31.3.11 else

Definition at line 1300 of file model-meat.hpp.

### 9.31.3.12 force\_new

```
bool force_new

Initial value:
{
    counter_fun.reset_array(&Array_)
```

Definition at line 428 of file model-meat.hpp.

#### 9.31.3.13 fun

```
Data_Counter_Type fun_
Initial value:
{
    counters->add_hash(fun_)
```

Definition at line 352 of file model-meat.hpp.

#### 9.31.3.14 i

```
const std::vector< double > size_t i = locator->second
```

Definition at line 1186 of file model-meat.hpp.

#### 9.31.3.15 i\_matches

```
int i_matches = -1
```

Definition at line 1306 of file model-meat.hpp.

#### 9.31.3.16 init\_fun\_

```
Data_Counter_Type Counter_fun_type<Array_Type,Data_Counter_Type> init_fun_
```

Definition at line 320 of file model-meat.hpp.

### 9.31.3.17 j

```
const std::vector< double > size_t size_t j = 0u
```

Definition at line 1288 of file model-meat.hpp.

### 9.31.3.18 k

```
size_t k = params.size()
```

Definition at line 1285 of file model-meat.hpp.

#### 9.31.3.19 key

```
std::vector< double > key = counters->gen_hash(Array_)
```

Definition at line 447 of file model-meat.hpp.

#### 9.31.3.20 locator

```
MapVec_type< double, size_t >::const_iterator locator = keys2support.find(key)
```

Definition at line 448 of file model-meat.hpp.

# 9.31.3.21 params

```
const std::vector< double > & params

Initial value:
{
    if (!this->with_pset)
        throw std::logic_error("Sampling is only available when store_pset() is active.")
```

Definition at line 1179 of file model-meat.hpp.

#### 9.31.3.22 probs

```
std::vector< double >& probs = pset_probs[a]
```

Definition at line 1289 of file model-meat.hpp.

#### 9.31.3.23 pset\_arrays

```
return this pset_arrays[a][j]
```

Definition at line 1340 of file model-meat.hpp.

#### 9.31.3.24 r

```
double r = urand(*rengine)
```

Definition at line 1282 of file model-meat.hpp.

#### 9.31.3.25 return

return

Definition at line 315 of file model-meat.hpp.

# 9.31.3.26 rule\_fun\_

```
Data_Rule_Dyn_Type rule_fun_
```

Definition at line 396 of file model-meat.hpp.

#### 9.31.3.27 rules

```
this rules
Initial value:
{
    rules->add_rule(rules, Data_Rule_Type())
```

Definition at line 362 of file model-meat.hpp.

### 9.31.3.28 rules\_

```
Data_Rule_Dyn_Type * rules_
Initial value:
{
    if (delete_rules)
        delete rules
```

Definition at line 371 of file model-meat.hpp.

#### 9.31.3.29 rules\_dyn

```
this rules_dyn = rules_
```

Definition at line 416 of file model-meat.hpp.

#### 9.31.3.30 stats

```
const std::vector< double >& stats = pset_stats[a]
```

Definition at line 1304 of file model-meat.hpp.

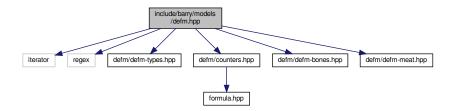
#### 9.31.3.31 stats\_support\_n\_arrays

```
stats_support_n_arrays[locator->second]
```

Definition at line 552 of file model-meat.hpp.

# 9.32 include/barry/models/defm.hpp File Reference

```
#include <iterator>
#include <regex>
#include "defm/defm-types.hpp"
#include "defm/counters.hpp"
#include "defm/defm-bones.hpp"
#include "defm/defm-meat.hpp"
Include dependency graph for defm.hpp:
```

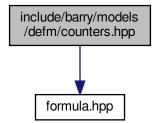


### **Namespaces**

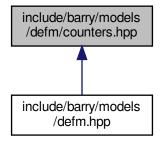
• defm

# 9.33 include/barry/models/defm/counters.hpp File Reference

```
#include "formula.hpp"
Include dependency graph for counters.hpp:
```



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



#### **Macros**

- #define MAKE DEFM HASHER(hasher, a, cov)
- #define DEFM\_RULEDYN\_LAMBDA(a)
- #define UNI SUB(a)

#### Macros for defining counters

- #define DEFM\_COUNTER(a) inline double (a) (const DEFMArray & Array, size\_t i, size\_t j, DEFMCounterData & data)
- #define DEFM\_COUNTER\_LAMBDA(a)

#### Macros for defining rules

- #define DEFM\_RULE(a) inline bool (a) (const DEFMArray & Array, size\_t i, size\_t j, bool & data)
- #define DEFM\_RULE\_LAMBDA(a)

#### **Functions**

void counter\_ones (DEFMCounters \*counters, int covar\_index=-1, std::string vname="", const std::vector< std::string > \*x\_names=nullptr)

Prevalence of ones.

- void counter\_logit\_intercept (DEFMCounters \*counters, size\_t n\_y, std::vector< size\_t > which={}, int covar\_index=-1, std::string vname="", const std::vector< std::string > \*x\_names=nullptr, const std::vector< std::string > \*y\_names=nullptr)

Prevalence of ones.

void counter\_transition\_formula (DEFMCounters \*counters, std::string formula, size\_t m\_order, size\_t n\_y, int covar\_index=-1, std::string vname="", const std::vector< std::string > \*x\_names=nullptr, const std::vector< std::string > \*y\_names=nullptr)

Prevalence of ones.

• void counter\_fixed\_effect (DEFMCounters \*counters, int covar\_index, double k, std::string vname="", const std::vector< std::string > \*x\_names=nullptr)

Prevalence of ones.

### Returns true if the cell is free

#### **Parameters**

```
rules | A pointer to a DEFMRules object (Rules < DEFMArray, bool > ).
```

- void rules\_markov\_fixed (DEFMRules \*rules, size\_t markov\_order)
   Number of edges.
- void rules\_dont\_become\_zero (DEFMSupport \*support, std::vector < size\_t > ids)
   Blocks switching a one to zero.

#### 9.33.1 Macro Definition Documentation

# 9.33.1.1 DEFM\_COUNTER

Function for definition of a network counter function

Definition at line 39 of file counters.hpp.

# 9.33.1.2 DEFM\_COUNTER\_LAMBDA

#### Value:

```
barry::Counter_fun_type<DEFMArray, DEFMCounterData> a = \
[](const DEFMArray & Array, size_t i, size_t j, DEFMCounterData & data) -> double
```

Lambda function for definition of a network counter function

Definition at line 43 of file counters.hpp.

## 9.33.1.3 DEFM\_RULE

Function for definition of a network counter function

Definition at line 53 of file counters.hpp.

# 9.33.1.4 DEFM\_RULE\_LAMBDA

#### Value:

```
barry::Rule_fun_type<DEFMArray, DEFMRuleData> a = \
[](const DEFMArray & Array, size_t i, size_t j, DEFMRuleData & data) -> bool
```

Lambda function for definition of a network counter function

Definition at line 57 of file counters.hpp.

# 9.33.1.5 DEFM\_RULEDYN\_LAMBDA

```
#define DEFM_RULEDYN_LAMBDA( a )
```

#### Value:

```
barry::Rule_fun_type<DEFMArray, DEFMRuleDynData> a = \
[](const DEFMArray & Array, size_t i, size_t j, DEFMRuleDynData & data) -> bool
```

Lambda function for definition of a network counter function

Definition at line 63 of file counters.hpp.

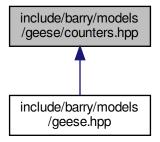
# 9.33.1.6 UNI\_SUB

## Value:

```
(\( (a) == 0) ? "\u2080" : (\( (a) == 1) ? "\u2081" : (\( (a) == 2) ? "\u2082" : (\( (a) == 3) ? "\u2083" : (\( (a) == 4) ? "\u2084" : (\( (a) == 5) ? "\u2085" : (\( (a) == 6) ? "\u2086" : (\( (a) == 7) ? "\u2087" : (\( (a) == 8) ? "\u2088" : \\ "\u2089"))))))))))))))
```

# 9.34 include/barry/models/geese/counters.hpp File Reference

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



## **Functions**

void rule\_dyn\_limit\_changes (PhyloSupport \*support, size\_t pos, size\_t lb, size\_t ub, size\_
 t duplication=Geese::etype default)

Overall functional gains.

- #define MAKE DUPL VARS()
- #define IS\_EITHER() (DATA\_AT == Geese::etype\_either)
- #define IS\_DUPLICATION() ((DATA\_AT == Geese::etype\_duplication) & (DPL))
- #define IS\_SPECIATION() ((DATA\_AT == Geese::etype\_speciation) & (!DPL))
- #define IF\_MATCHES()
- #define IF\_NOTMATCHES()
- #define PHYLO COUNTER LAMBDA(a)

Extension of a simple counter.

- #define PHYLO\_RULE\_DYN\_LAMBDA(a)
- #define PHYLO CHECK MISSING()
- std::string get\_last\_name (size\_t d)
- void counter\_overall\_gains (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

Overall functional gains.

- void counter\_gains (PhyloCounters \*counters, std::vector < size\_t > nfun, size\_t duplication=Geese::etype\_default)

  Functional gains for a specific function (nfun).
- void counter\_gains\_k\_offspring (PhyloCounters \*counters, std::vector< size\_t > nfun, size\_t k=1u, size\_t duplication=Geese::etype\_default)

k genes gain function nfun

- · void counter\_genes\_changing (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)
  - Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)
- void counter\_preserve\_pseudogene (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_ 
  t duplication=Geese::etype\_default)

Keeps track of how many pairs of genes preserve pseudostate.

· void counter prop genes changing (PhyloCounters \*counters, size t duplication=Geese::etype default)

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

- void counter\_overall\_loss (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

  Overall functional loss.
- void counter\_maxfuns (PhyloCounters \*counters, size\_t lb, size\_t ub, size\_t duplication=Geese::etype\_default)

  Cap the number of functions per gene.
- void counter\_loss (PhyloCounters \*counters, std::vector < size\_t > nfun, size\_t duplication=Geese::etype\_default)

  Total count of losses for an specific function.
- void counter\_overall\_changes (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

  Total number of changes. Use this statistic to account for "preservation".
- void counter\_subfun (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

  Total count of Sub-functionalization events.
- void counter\_cogain (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

  Co-evolution (joint gain or loss)
- void counter\_longest (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

  Longest branch mutates (either by gain or by loss)
- void counter\_neofun (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

  Total number of neofunctionalization events.
- void counter\_pairwise\_neofun\_singlefun (PhyloCounters \*counters, size\_t nfunA, size\_t duplication=Geese::etype\_default)

  Total number of neofunctionalization events sum\_u sum\_{{w < u} [x(u,a)\*(1 x(w,a)) + (1 x(u,a)) \* x(w,a)] change stat: delta{x(u,a): 0->1} = 1 2 \* x(w,a)
- void counter\_neofun\_a2b (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

  Total number of neofunctionalization events.
- void counter\_co\_opt (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default) Function co-opting.
- void counter\_k\_genes\_changing (PhyloCounters \*counters, size\_t k, size\_t duplication=Geese::etype\_default)

  Indicator function. Equals to one if k genes changed and zero otherwise.
- void counter\_less\_than\_p\_prop\_genes\_changing (PhyloCounters \*counters, double p, size\_t duplication=Geese::etype\_default
- void counter\_gains\_from\_0 (PhyloCounters \*counters, std::vector < size\_t > nfun, size\_t duplication=Geese::etype\_default)

  Used when all the functions are in 0 (like the root node prob.)
- void counter\_overall\_gains\_from\_0 (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

  Used when all the functions are in 0 (like the root node prob.)
- void counter\_pairwise\_overall\_change (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

  Used when all the functions are in 0 (like the root node prob.)
- void counter\_pairwise\_preserving (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_
   t duplication=Geese::etype\_default)

Used when all the functions are in 0 (like the root node prob.)

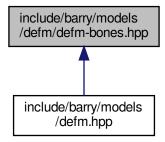
Indicator function. Equals to one if k genes changed and zero otherwise.

void counter\_pairwise\_first\_gain (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_
 t duplication=Geese::etype\_default)

Used when all the functions are in 0 (like the root node prob.)

# 9.35 include/barry/models/defm/defm-bones.hpp File Reference

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

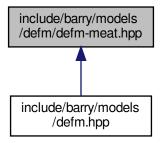


## **Classes**

· class DEFM

# 9.36 include/barry/models/defm/defm-meat.hpp File Reference

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



#### **Macros**

- #define DEFM\_RANGES(a)
- #define DEFM\_LOOP\_ARRAYS(a) for (size\_t a = 0u; a < (nobs\_i M\_order); ++a)

## **Functions**

• std::vector< double > keygen\_defm (const DEFMArray &Array\_, DEFMCounterData \*data)

# 9.36.1 Macro Definition Documentation

# 9.36.1.1 DEFM\_LOOP\_ARRAYS

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

# 9.36.1.2 DEFM\_RANGES

Definition at line 30 of file defm-meat.hpp.

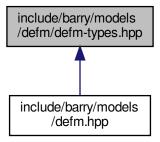
# 9.36.2 Function Documentation

#### 9.36.2.1 keygen\_defm()

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

# 9.37 include/barry/models/defm/defm-types.hpp File Reference

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



## **Classes**

- class DEFMData
  - Data class for DEFM arrays.
- class DEFMCounterData

Data class used to store arbitrary size\_t or double vectors.

- class DEFMRuleData
- · class DEFMRuleDynData

# **Typedefs**

• typedef barry::BArrayDense< int, DEFMData > DEFMArray

# Convenient typedefs for network objects.

- typedef barry::Counter< DEFMArray, DEFMCounterData > DEFMCounter
- typedef barry::Counters < DEFMArray, DEFMCounterData > DEFMCounters
- typedef barry::Support
   DEFMArray, DEFMCounterData, DEFMRuleData, DEFMRuleDynData > DEFMSupport
- typedef barry::StatsCounter< DEFMArray, DEFMCounterData > DEFMStatsCounter
- typedef barry::Model DEFMArray, DEFMCounterData, DEFMRuleData, DEFMRuleDynData > DEFMModel
- typedef barry::Rule < DEFMArray, DEFMRuleData > DEFMRule
- typedef barry::Rules< DEFMArray, DEFMRuleData > DEFMRules
- typedef barry::Rule < DEFMArray, DEFMRuleDynData > DEFMRuleDyn
- typedef barry::Rules < DEFMArray, DEFMRuleDynData > DEFMRulesDyn

# 9.37.1 Typedef Documentation

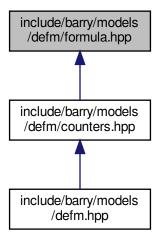
# 9.37.1.1 **DEFMArray**

typedef barry::BArrayDense<int, DEFMData> DEFMArray

Definition at line 5 of file defm-types.hpp.

# 9.38 include/barry/models/defm/formula.hpp File Reference

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



# **Functions**

void defm\_motif\_parser (std::string formula, std::vector< size\_t > &locations, std::vector< bool > &signs, size t m order, size t y ncol)

Parses a motif formula.

# 9.38.1 Function Documentation

#### 9.38.1.1 defm\_motif\_parser()

Parses a motif formula.

This function will take the formula and generate the corresponding input for defm::counter\_transition(). Formulas can be specified in the following ways:

- Intercept effect: {...} No transition, only including the current state.
- Transition effect: {...} > {...} Includes current and previous states.

The general notation is  $[0]y[column id]_[row id]$ . A preceding zero means that the value of the cell is considered to be zero. The column id goes between 0 and the number of columns in the array - 1 (so it is indexed from 0,) and the row id goes from 0 to m\_order.

#### Intercept effects

Intercept effects only involve a single set of curly brackets. Using the 'greater-than' symbol (i.e., '<') is only for transition effects. When specifying intercept effects, users can skip the  $row_id$ , e.g.,  $y0_0$  is equivalent to y0. If the passed  $row_id$  is different from the Markov order, i.e.,  $row_id$  !=  $m_order$ , then the function returns with an error.

## Examples:

• " $\{y0, 0y1\}$ " is equivalent to set a motif with the first element equal to one and the second to zero.

# **Transition effects**

Transition effects can be specified using two sets of curly brackets and an greater-than symbol, i.e.,  $\{\ldots\}$  >  $\{\ldots\}$ . The first set of brackets, which we call LHS, can only hold row id that are less than m\_order.

#### **Parameters**

formula	
locations	
signs	
m_order	
y_ncol	

Definition at line 46 of file formula.hpp.

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

```
#include "geese/geese-types.hpp"
#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/geese-meat-predict_exhaust.hpp"
#include "geese/geese-meat-predict_sim.hpp"
#include "geese/flock-bones.hpp"
#include "geese/flock-meat.hpp"
#include "geese/counters.hpp"
#include dependency graph for geese.hpp:
```

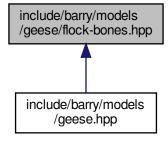


# **Namespaces**

• geese

# 9.40 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.

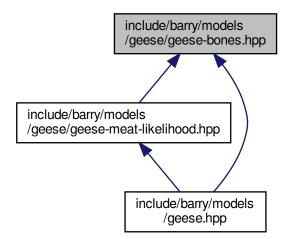
# 9.41 include/barry/models/geese/flock-meat.hpp File Reference

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



# 9.42 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 PhyloArray &array, const PhyloCounterData *d)
```

bool vec diff (const std::vector< size t > &s, const std::vector< size t > &a)

# 9.42.1 Macro Definition Documentation

## **9.42.1.1 INITIALIZED**

```
#define INITIALIZED( )

Value:
    if (!this->initialized) \
    throw std::logic_error("The model has not been initialized yet.");
```

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

#### 9.42.2 Function Documentation

# 9.42.2.1 keygen\_full()

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

#### 9.42.2.2 RULE\_FUNCTION()

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

# 9.42.2.3 vec\_diff()

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

# 9.42.2.4 vector\_caster()

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

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



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

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



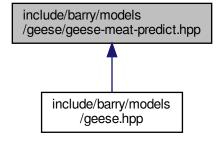


# 9.45 include/barry/models/geese/geese-meat-likelihood\_exhaust.hpp File Reference

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

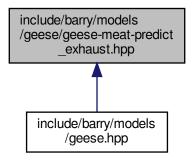


# 9.46 include/barry/models/geese/geese-meat-predict.hpp File Reference

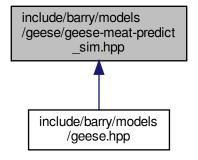


# 9.47 include/barry/models/geese/geese-meat-predict\_exhaust.hpp File Reference

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



# 9.48 include/barry/models/geese/geese-meat-predict\_sim.hpp File Reference

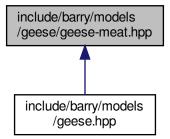


# 9.49 include/barry/models/geese/geese-meat-simulate.hpp File Reference

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

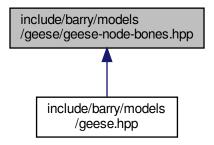


# 9.50 include/barry/models/geese/geese-meat.hpp File Reference



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

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



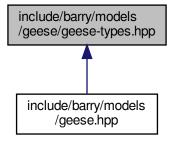
# Classes

class Node

A single node for the model.

# 9.52 include/barry/models/geese/geese-types.hpp File Reference

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



## **Classes**

class NodeData

Data definition for the PhyloArray class.

- class PhyloCounterData
- class PhyloRuleDynData

# **Typedefs**

#### Convenient typedefs for Node objects.

- typedef std::vector< std::pair< size t, size t >> PhyloRuleData
- typedef barry::BArrayDense< size\_t, NodeData > PhyloArray
- typedef barry::Counter< PhyloArray, PhyloCounterData > PhyloCounter
- typedef barry::Counters < PhyloArray, PhyloCounterData > PhyloCounters
- typedef barry::Rule < PhyloArray, PhyloRuleData > PhyloRule
- typedef barry::Rules
   PhyloArray, PhyloRuleData
   PhyloRules
- typedef barry::Rule < PhyloArray, PhyloRuleDynData > PhyloRuleDyn
- typedef barry::Rules
   PhyloArray, PhyloRuleDynData
   PhyloRulesDyn
- typedef barry::Support
   PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloSupport
- typedef barry::StatsCounter< PhyloArray, PhyloCounterData > PhyloStatsCounter
- typedef barry::Model < PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloModel
- typedef barry::PowerSet
   PhyloArray, PhyloRuleData > PhyloPowerSet

# 9.52.1 Typedef Documentation

#### 9.52.1.1 PhyloArray

typedef barry::BArrayDense<size\_t, NodeData> PhyloArray

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

#### 9.52.1.2 PhyloCounter

typedef barry::Counter<PhyloArray, PhyloCounterData > PhyloCounter

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

#### 9.52.1.3 PhyloCounters

typedef barry::Counters< PhyloArray, PhyloCounterData> PhyloCounters

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

#### 9.52.1.4 PhyloModel

typedef barry::Model<PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloModel

Definition at line 113 of file geese-types.hpp.

# 9.52.1.5 PhyloPowerSet

typedef barry::PowerSet<PhyloArray, PhyloRuleData> PhyloPowerSet

Definition at line 114 of file geese-types.hpp.

## 9.52.1.6 PhyloRule

typedef barry::Rule<PhyloArray,PhyloRuleData> PhyloRule

Definition at line 105 of file geese-types.hpp.

## 9.52.1.7 PhyloRuleData

typedef std::vector< std::pair< size\_t, size\_t > > PhyloRuleData

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

# 9.52.1.8 PhyloRuleDyn

typedef barry::Rule<PhyloArray,PhyloRuleDynData> PhyloRuleDyn

Definition at line 108 of file geese-types.hpp.

# 9.52.1.9 PhyloRules

typedef barry::Rules<PhyloArray,PhyloRuleData> PhyloRules

Definition at line 106 of file geese-types.hpp.

# 9.52.1.10 PhyloRulesDyn

typedef barry::Rules<PhyloArray,PhyloRuleDynData> PhyloRulesDyn

Definition at line 109 of file geese-types.hpp.

# 9.52.1.11 PhyloStatsCounter

typedef barry::StatsCounter<PhyloArray, PhyloCounterData> PhyloStatsCounter

Definition at line 112 of file geese-types.hpp.

# 9.52.1.12 PhyloSupport

typedef barry::Support<PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloSupport

Definition at line 111 of file geese-types.hpp.

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

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



# **Classes**

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

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



# 9.55 include/barry/progress.hpp File Reference

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



# **Classes**

• class Progress

A simple progress bar.

# **Macros**

• #define BARRY\_PROGRESS\_BAR\_WIDTH 80

#### 9.55.1 Macro Definition Documentation

## 9.55.1.1 BARRY PROGRESS BAR WIDTH

```
#define BARRY_PROGRESS_BAR_WIDTH 80
```

Definition at line 5 of file progress.hpp.

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

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



# **Classes**

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

# **Functions**

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

# 9.56.1 Function Documentation

# 9.56.1.1 rule\_fun\_default()

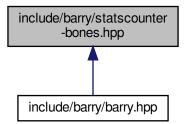
Definition at line 5 of file rules-bones.hpp.

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

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



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



#### Classes

 $\bullet \ \ {\it class StatsCounter} < {\it Array\_Type}, \ {\it Data\_Type} >$ 

Count stats for a single Array.

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

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



# **Macros**

- #define STATSCOUNTER TYPE() StatsCounter<Array Type, Data Type>
- #define STATSCOUNTER\_TEMPLATE\_ARGS() < typename Array\_Type, typename Data\_Type >
- #define STATSCOUNTER\_TEMPLATE(a, b) template STATSCOUNTER\_TEMPLATE\_ARGS() inline a STATSCOUNTER\_TYPE()::b

# **Functions**

- STATSCOUNTER\_TEMPLATE (, StatsCounter)(const StatsCounter < Array\_Type
- EmptyArray clear ()
- STATSCOUNTER TEMPLATE (,~StatsCounter)()
- STATSCOUNTER\_TEMPLATE (void, reset\_array)(const Array\_Type \*Array\_)
- STATSCOUNTER\_TEMPLATE (void, add\_counter)(Counter< Array\_Type</li>
- STATSCOUNTER\_TEMPLATE (void, set\_counters)(Counters< Array\_Type
- STATSCOUNTER\_TEMPLATE (void, count\_init)(size\_t i
- current\_stats resize (counters->size(), 0.0)
- for (size\_t n=0u;n< counters->size();++n) current\_stats[n]
- STATSCOUNTER TEMPLATE (void, count current)(size ti
- STATSCOUNTER\_TEMPLATE (std::vector< std::string >, get\_names)() const
- STATSCOUNTER\_TEMPLATE (std::vector< std::string >, get\_descriptions)() const

## **Variables**

size\_t j

```
    Data_Type & counter
    EmptyArray = *Array
    current_stats = counter.current_stats
    counters = new Counters<Array_Type,Data_Type>((*counter.counters))
    counter_deleted = false
    Data_Type f_
    return
    Data_Type * counters_
```

#### 9.59.1 Macro Definition Documentation

## 9.59.1.1 STATSCOUNTER\_TEMPLATE

Definition at line 8 of file statscounter-meat.hpp.

## 9.59.1.2 STATSCOUNTER\_TEMPLATE\_ARGS

```
template STATSCOUNTER_TEMPLATE_ARGS() <typename Array_Type, typename Data_Type>
```

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

#### 9.59.1.3 STATSCOUNTER TYPE

```
template Data_Type * STATSCOUNTER_TYPE( ) StatsCounter<Array_Type,Data_Type>
```

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

# 9.59.2 Function Documentation

# 9.59.2.1 clear()

```
EmptyArray clear ( )
```

# 9.59.2.2 for()

## 9.59.2.3 resize()

# 9.59.2.4 STATSCOUNTER\_TEMPLATE() [1/9]

```
STATSCOUNTER_TEMPLATE (
StatsCounter ) const
```

# 9.59.2.5 STATSCOUNTER\_TEMPLATE() [2/9]

```
STATSCOUNTER_TEMPLATE ( \sim \textit{StatsCounter} \ )
```

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

# 9.59.2.6 STATSCOUNTER\_TEMPLATE() [3/9]

```
STATSCOUNTER_TEMPLATE (
          std::vector< std::string > ,
          get_descriptions ) const
```

Definition at line 256 of file statscounter-meat.hpp.

# 9.59.2.7 STATSCOUNTER\_TEMPLATE() [4/9]

```
STATSCOUNTER_TEMPLATE (
          std::vector< std::string > ,
          get_names ) const
```

Definition at line 251 of file statscounter-meat.hpp.

# 9.59.2.8 STATSCOUNTER\_TEMPLATE() [5/9]

# 9.59.2.9 STATSCOUNTER\_TEMPLATE() [6/9]

# 9.59.2.10 STATSCOUNTER\_TEMPLATE() [7/9]

# 9.59.2.11 STATSCOUNTER\_TEMPLATE() [8/9]

```
STATSCOUNTER_TEMPLATE (

void ,

reset_array ) const
```

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

# 9.59.2.12 STATSCOUNTER\_TEMPLATE() [9/9]

# 9.59.3 Variable Documentation

## 9.59.3.1 counter

```
Data_Type& counter

Initial value:
{
    Array = counter.Array
```

Definition at line 12 of file statscounter-meat.hpp.

# 9.59.3.2 counter\_deleted

```
counter_deleted = false
```

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

## 9.59.3.3 counters

```
counters = new Counters<Array_Type,Data_Type>((*counter.counters))
```

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

#### 9.59.3.4 counters\_

```
Data_Type* counters_
Initial value:
{
    if (!counter_deleted)
        delete counters
```

Definition at line 53 of file statscounter-meat.hpp.

# 9.59.3.5 current\_stats

```
current_stats = counter.current_stats
```

Definition at line 19 of file statscounter-meat.hpp.

## 9.59.3.6 EmptyArray

```
EmptyArray = *Array
```

Definition at line 17 of file statscounter-meat.hpp.

```
9.59.3.7 f_
```

```
Data_Rule_Dyn_Type f_
Initial value:
{
    counters->add_counter(f_)
```

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

```
9.59.3.8 j
```

```
size_t j
```

# Initial value:

```
if (counters->size() == 0u)
    throw std::logic_error("No counters added: Cannot count without knowning what to count!")
```

Definition at line 66 of file statscounter-meat.hpp.

#### 9.59.3.9 return

return

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

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

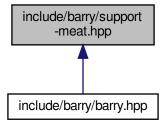


#### Classes

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

# 9.61 include/barry/support-meat.hpp File Reference

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



#### **Macros**

- #define BARRY\_SUPPORT\_MEAT\_HPP 1
- #define SUPPORT\_TEMPLATE\_ARGS()
- #define SUPPORT\_TYPE()
- #define SUPPORT\_TEMPLATE(a, b)

# **Functions**

- SUPPORT\_TEMPLATE (void, init\_support)(std
- SUPPORT\_TEMPLATE (void, reset\_array)()
- SUPPORT\_TEMPLATE (void, reset\_array)(const Array\_Type &Array\_)
- SUPPORT\_TEMPLATE (void, calc\_backend\_sparse)(size\_t pos
- · calc backend sparse (pos+1u, array bank, stats bank)
- EmptyArray insert\_cell (coord\_i, coord\_j, EmptyArray.default\_val().value, false, false)
- for (size\_t n=0u;n< n\_counters;++n)
- if (rules\_dyn->size() > 0u)
- if (array\_bank !=nullptr) array\_bank -> push\_back(EmptyArray)
- EmptyArray rm cell (coord i, coord j, false, false)
- if (change\_stats\_different > 0u)
- SUPPORT\_TEMPLATE (void, calc\_backend\_dense)(size\_t pos
- calc\_backend\_dense (pos+1u, array\_bank, stats\_bank)
- EmptyArray insert\_cell (coord\_i, coord\_j, 1, false, false)
- SUPPORT\_TEMPLATE (void, calc)(std
- SUPPORT\_TEMPLATE (void, add\_counter)(Counter< Array\_Type</li>
- SUPPORT\_TEMPLATE (void, set\_counters)(Counters < Array\_Type</li>

```
• SUPPORT_TEMPLATE (void, add_rule)(Rule < Array_Type
```

- SUPPORT\_TEMPLATE (void, set\_rules)(Rules< Array\_Type
- SUPPORT TEMPLATE (void, add rule dyn)(Rule < Array Type
- SUPPORT\_TEMPLATE (void, set\_rules\_dyn)(Rules< Array\_Type
- SUPPORT TEMPLATE (bool, eval rules dyn)(const std
- SUPPORT\_TEMPLATE (std::vector< double >, get\_counts)() const
- SUPPORT TEMPLATE (std::vector< double > \*, get current stats)()
- SUPPORT\_TEMPLATE (void, print)() const
- SUPPORT\_TEMPLATE (const FreqTable < double > &, get\_data)() const

#### **Variables**

- std::vector< Array\_Type > \* array\_bank
- std::vector< Array\_Type > std::vector< double > \* stats\_bank
- const size\_t & coord\_i = coordinates\_free[pos \* 2u]
- const size t & coord j = coordinates free[pos \* 2u + 1u]
- · double tmp chng
- size\_t change\_stats\_different = hashes\_initialized[pos] ? 0u : 1u
- else
- & hashes [pos]
- return
- Data\_Counter\_Type f\_
- Data Counter Type \* counters
- delete counters = false
- counters = counters
- Data\_Rule\_Type \* rules\_
- delete\_rules = false
- rules = rules
- delete\_rules\_dyn = false
- rules\_dyn = rules\_

#### 9.61.1 Macro Definition Documentation

# 9.61.1.1 BARRY\_SUPPORT\_MEAT\_HPP

```
#define BARRY_SUPPORT_MEAT_HPP 1
```

Definition at line 2 of file support-meat.hpp.

## 9.61.1.2 SUPPORT\_TEMPLATE

#### Value:

```
template SUPPORT_TEMPLATE_ARGS() \
inline a SUPPORT_TYPE()::b
```

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

# 9.61.1.3 SUPPORT\_TEMPLATE\_ARGS

```
Value:
    <typename Array_Type, typename \
    Data_Counter_Type, typename Data_Rule_Type, typename Data_Rule_Dyn_Type>
```

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

# 9.61.1.4 SUPPORT\_TYPE

```
template Data_Rule_Dyn_Type * SUPPORT_TYPE()

Value:
    Support<Array_Type, Data_Counter_Type, Data_Rule_Type,\
    Data_Rule_Dyn_Type>
```

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

# 9.61.2 Function Documentation

# 9.61.2.1 calc\_backend\_dense()

```
calc_backend_dense (
    pos+ 1u,
    array_bank ,
    stats_bank )
```

# 9.61.2.2 calc\_backend\_sparse()

```
calc_backend_sparse (
    pos+ 1u,
    array_bank ,
    stats_bank )
```

## 9.61.2.3 for()

```
for ( )
```

Definition at line 159 of file support-meat.hpp.

# **9.61.2.4** if() [1/3]

# 9.61.2.5 if() [2/3]

```
if ( \label{eq:change_stats_different} \mbox{,} \\ \mbox{Ou })
```

Definition at line 239 of file support-meat.hpp.

## 9.61.2.6 if() [3/3]

```
if (
    rules_dyn-> size(),
    0u )
```

Definition at line 187 of file support-meat.hpp.

# 9.61.2.7 insert\_cell() [1/2]

# 9.61.2.8 insert\_cell() [2/2]

# 9.61.2.9 rm\_cell()

#### 9.61.2.10 SUPPORT\_TEMPLATE() [1/17]

```
SUPPORT_TEMPLATE (
          bool ,
          eval_rules_dyn ) const
```

Definition at line 489 of file support-meat.hpp.

# 9.61.2.11 SUPPORT\_TEMPLATE() [2/17]

Definition at line 558 of file support-meat.hpp.

# 9.61.2.12 SUPPORT\_TEMPLATE() [3/17]

```
SUPPORT_TEMPLATE (
          std::vector< double > * ,
          get_current_stats )
```

Definition at line 543 of file support-meat.hpp.

# 9.61.2.13 SUPPORT\_TEMPLATE() [4/17]

```
SUPPORT_TEMPLATE (
          std::vector< double > ,
          get_counts ) const
```

Definition at line 531 of file support-meat.hpp.

# 9.61.2.14 SUPPORT\_TEMPLATE() [5/17]

```
SUPPORT_TEMPLATE (
     void ,
     add_counter )
```

# 9.61.2.15 SUPPORT\_TEMPLATE() [6/17]

# 9.61.2.16 SUPPORT\_TEMPLATE() [7/17]

## 9.61.2.17 SUPPORT\_TEMPLATE() [8/17]

```
SUPPORT_TEMPLATE (
    void ,
    calc )
```

Definition at line 367 of file support-meat.hpp.

# 9.61.2.18 SUPPORT\_TEMPLATE() [9/17]

# 9.61.2.19 SUPPORT\_TEMPLATE() [10/17]

## 9.61.2.20 SUPPORT\_TEMPLATE() [11/17]

Definition at line 13 of file support-meat.hpp.

## 9.61.2.21 SUPPORT\_TEMPLATE() [12/17]

```
SUPPORT_TEMPLATE (
     void ,
     print ) const
```

Definition at line 547 of file support-meat.hpp.

#### 9.61.2.22 SUPPORT\_TEMPLATE() [13/17]

```
SUPPORT_TEMPLATE (
     void ,
     reset_array )
```

Definition at line 114 of file support-meat.hpp.

#### 9.61.2.23 SUPPORT\_TEMPLATE() [14/17]

Definition at line 120 of file support-meat.hpp.

## 9.61.2.24 SUPPORT\_TEMPLATE() [15/17]

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## 9.61.2.25 SUPPORT\_TEMPLATE() [16/17]

## 9.61.2.26 SUPPORT\_TEMPLATE() [17/17]

#### 9.61.3 Variable Documentation

## 9.61.3.1 array\_bank

```
\verb|std::vector< Array_Type| > * array_bank|
```

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

#### 9.61.3.2 change\_stats\_different

```
size_t change_stats_different = hashes_initialized[pos] ? Ou : 1u
```

Definition at line 158 of file support-meat.hpp.

#### 9.61.3.3 coord i

```
const size_t & coord_i = coordinates_free[pos * 2u]
```

Definition at line 144 of file support-meat.hpp.

## 9.61.3.4 coord\_j

```
const size_t & coord_j = coordinates_free[pos * 2u + 1u]
```

Definition at line 145 of file support-meat.hpp.

#### 9.61.3.5 counters

```
counters = counters_
```

Definition at line 417 of file support-meat.hpp.

#### 9.61.3.6 counters\_

```
Data_Counter_Type* counters_
```

#### Initial value:

{

```
if (delete_counters)
    delete counters
```

Definition at line 410 of file support-meat.hpp.

#### 9.61.3.7 delete\_counters

```
delete_counters = false
```

Definition at line 416 of file support-meat.hpp.

## 9.61.3.8 delete\_rules

```
delete_rules = false
```

Definition at line 450 of file support-meat.hpp.

## 9.61.3.9 delete\_rules\_dyn

```
delete_rules_dyn = false
```

Definition at line 482 of file support-meat.hpp.

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#### 9.61.3.10 else

Definition at line 212 of file support-meat.hpp.

```
9.61.3.11 f_
```

```
Data_Rule_Dyn_Type f_
```

## Initial value:

```
counters->add_counter(f_)
```

Definition at line 401 of file support-meat.hpp.

#### 9.61.3.12 hashes

& hashes

Definition at line 217 of file support-meat.hpp.

#### 9.61.3.13 return

return

Definition at line 249 of file support-meat.hpp.

## 9.61.3.14 rules

```
rules = rules_
```

Definition at line 451 of file support-meat.hpp.

## 9.61.3.15 rules\_

```
Data_Rule_Dyn_Type* rules_
Initial value:
{
    if (delete_rules)
        delete rules
```

Definition at line 444 of file support-meat.hpp.

## 9.61.3.16 rules\_dyn

```
rules_dyn = rules_
```

Definition at line 483 of file support-meat.hpp.

#### 9.61.3.17 stats\_bank

```
std::vector< Array_Type > std::vector< double > * stats_bank

Initial value:
{
    if (pos >= coordiantes_n_free)
```

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

## 9.61.3.18 tmp\_chng

```
double tmp_chng
```

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

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# 9.62 include/barry/typedefs.hpp File Reference

```
#include "barry-configuration.hpp"
#include "barry-debug.hpp"
#include "progress.hpp"
Include dependency graph for typedefs.hpp:
```

barry-configuration.hpp barry-debug.hpp progress.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.

## **Typedefs**

```
    typedef std::vector< std::pair< std::vector< double >, size t >> Counts type

    template<typename Cell_Type >
      using Row_type = Map< size_t, Cell< Cell_Type > >
    template<typename Cell_Type >
      using Col_type = Map< size_t, Cell< Cell_Type > * >
    • template<typename Ta = double, typename Tb = size_t>
      using MapVec_type = std::unordered_map< std::vector< Ta >, Tb, vecHasher< Ta >>
    • template<typename Array_Type , typename Data_Type >
      using Hasher_fun_type = std::function < std::vector < double >(const Array_Type &, Data_Type *)>
          Hasher function used by the counter.

    template<typename Array_Type , typename Data_Type >

      using Counter_fun_type = std::function< double(const Array_Type &, size_t, size_t, Data_Type &)>
          Counter and rule functions.
    • template<typename Array_Type , typename Data_Type >
      using Rule_fun_type = std::function< bool(const Array_Type &, size_t, Data_Type &)>
Functions

    std::vector < size_t > sort_array (const double *v, size_t start, size_t ncols, size_t nrows)

          Ascending sorting an array.
    • template<typename T >
      T vec_inner_prod (const T *a, const T *b, size_t n)
    • template<> double vec_inner_prod (const double *a, const double *b, size_t n)
```

```
template<typename T >
  bool vec equal (const std::vector< T > &a, const std::vector< T > &b)
     Compares if -a- and -b- are equal.
```

• template<typename T > bool vec\_equal\_approx (const std::vector< T > &a, const std::vector< T > &b, double eps=1e-100)

#### **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
```

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## 9.62.1 Typedef Documentation

#### 9.62.1.1 Col type

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

Definition at line 70 of file typedefs.hpp.

#### 9.62.1.2 Counter\_fun\_type

```
template<typename Array_Type , typename Data_Type >
using Counter_fun_type = std::function<double(const Array_Type &, size_t, size_t, Data_Type &)>
```

Counter and rule functions.

#### **Parameters**

Array_Type	a BArray
unit,size⇔	Focal cell
_t	
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 187 of file typedefs.hpp.

#### 9.62.1.3 Counts\_type

```
typedef std::vector< std::pair< std::vector<double>, size_t >> Counts_type
Definition at line 51 of file typedefs.hpp.
```

## 9.62.1.4 Hasher\_fun\_type

```
template<typename Array_Type , typename Data_Type >
using Hasher_fun_type = std::function<std::vector<double>(const Array_Type &, Data_Type *)>
```

Hasher function used by the counter.

Used to characterize the support of the array.

**Template Parameters** 

```
Array_Type
```

Definition at line 200 of file typedefs.hpp.

## 9.62.1.5 MapVec\_type

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

Definition at line 128 of file typedefs.hpp.

## 9.62.1.6 Row\_type

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

Definition at line 67 of file typedefs.hpp.

#### 9.62.1.7 Rule\_fun\_type

```
template<typename Array_Type , typename Data_Type >
using Rule_fun_type = std::function<bool(const Array_Type &, size_t, size_t, Data_Type &)>
```

Definition at line 190 of file typedefs.hpp.

## 9.62.2 Function Documentation

#### 9.62.2.1 sort\_array()

Ascending sorting an array.

It will sort an array solving ties using the next column. Data is stored column-wise.

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#### **Template Parameters**



#### **Parameters**



#### Returns

std::vector<size\_t> The sorting index.

Definition at line 141 of file typedefs.hpp.

## 9.62.2.2 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 210 of file typedefs.hpp.

#### 9.62.2.3 vec\_equal\_approx()

Definition at line 235 of file typedefs.hpp.

## 9.62.2.4 vec\_inner\_prod() [1/2]

Definition at line 286 of file typedefs.hpp.

## 9.62.2.5 vec\_inner\_prod() [2/2]

Definition at line 263 of file typedefs.hpp.

## 9.63 README.md File Reference

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