barry: Your go-to motif accountant 0.0-1

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Chapter 1

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."

2 Main Page

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 here and in the PDF version of the documentation here.

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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Chapter 2

Module Index

2.1 Modules

Here is a list of all modules:

Counting													 										13
Statistical Models																							24
Network counters																							24
Phylo counters																							49
Phylo rules												 						 					36

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Chapter 3

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 > , bool >
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 {>\ } $
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, bool >
vecHasher< T >

Chapter 4

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 117
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 143
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
Flock
A Flock is a group of Geese

10 Class Index

FreqTable < T >	
Frequency table of vectors	159
Geese	
Annotated Phylo Model	163
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:	177
NetCounterData	
Data class used to store arbitrary size_t or double vectors	205
NetworkData	
Data class for Networks	206
Node	
A single node for the model	209
NodeData	
Data definition for the PhyloArray class	215
PhyloCounterData	216
PhyloRuleDynData	219
PowerSet < Array_Type, Data_Rule_Type >	
Powerset of a binary array	222
Progress	
A simple progress bar	228
Rule < Array_Type, Data_Type >	
Rule for determining if a cell should be included in a sequence	229
Rules< Array_Type, Data_Type >	
Vector of objects of class Rule	232
StatsCounter< Array_Type, Data_Type >	
Count stats for a single Array	236
Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >	
Compute the support of sufficient statistics	240
vecHasher< T >	251

Chapter 5

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
$include/barry/models/defm.hpp \\ \dots \\$
include/barry/models/geese.hpp
include/barry/models/defm/counters.hpp
$include/barry/models/defm/defm-bones.hpp \\ \dots \dots$
include/barry/models/defm/defm-meat.hpp
$include/barry/models/defm/defm-types.hpp \\ \dots \dots$
include/barry/models/defm/formula.hpp
include/barry/models/geese/counters.hpp
$include/barry/models/geese/flock-bones.hpp \\ \dots \\$
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
$include/barry/models/geese/geese-meat-predict_sim.hpp \\ \dots \\$
include/barry/models/geese/geese-meat-simulate.hpp
include/barry/models/geese/geese-meat.hpp
$include/barry/models/geese/geese-node-bones.hpp \\ \dots \\$
include/barry/models/geese/geese-types.hpp

Chapter 6

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)

Data for the counters.

- #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_RULE_LAMBDA(a)

Extension of a simple counter.

- #define PHYLO_COUNTER_LAMBDA(a)
- #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.)

6.1 Counting 15

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

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

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:

```
barry::Hasher_fun_type<DEFMArray, DEFMCounterData>
  hasher = [cov](const DEFMArray & array, DEFMCounterData * d) -> 
  std::vector< double > {
    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>
```

Data for the counters.

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

This class is used to store the data for the counters. It is used by the ${\tt Counters}$ class.

Definition at line 27 of file counters.hpp.

6.1.2.7 MAKE_DUPL_VARS

```
#define MAKE_DUPL_VARS( )

Value:
    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.

6.1 Counting 17

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 45 of file counters.hpp.

6.1.2.9 PHYLO_COUNTER_LAMBDA

Definition at line 39 of file counters.hpp.

6.1.2.10 PHYLO_RULE_DYN_LAMBDA

Definition at line 42 of file counters.hpp.

6.1.2.11 PHYLO RULE 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.

6.1.3 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 1299 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 794 of file counters.hpp.

6.1.3.3 counter_gains()

Functional gains for a specific function (nfun).

Definition at line 99 of file counters.hpp.

6.1 Counting 19

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 1633 of file counters.hpp.

6.1.3.5 counter_gains_k_offspring()

k genes gain function nfun

Definition at line 159 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 231 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 1397 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 1517 of file counters.hpp.

6.1.3.9 counter_longest()

Longest branch mutates (either by gain or by loss)

Definition at line 851 of file counters.hpp.

6.1.3.10 counter_loss()

Total count of losses for an specific function.

Definition at line 594 of file counters.hpp.

6.1.3.11 counter_maxfuns()

Cap the number of functions per gene.

Definition at line 532 of file counters.hpp.

6.1 Counting 21

6.1.3.12 counter_neofun()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1021 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 1166 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 646 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 61 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 1699 of file counters.hpp.

6.1.3.17 counter_overall_loss()

Overall functional loss.

Definition at line 484 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 1951 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 1102 of file counters.hpp.

6.1 Counting 23

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 1747 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 1812 of file counters.hpp.

6.1.3.22 counter_preserve_pseudogene()

Keeps track of how many pairs of genes preserve pseudostate.

Definition at line 300 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 382 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 705 of file counters.hpp.

6.1.3.25 get_last_name()

Definition at line 48 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.

6.3 Network counters 25

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.

• void rule_constrain_support (DEFMSupport *support, size_t pos, double lb, double ub)

Overall functional gains.

6.3.1 Detailed Description

Counters for network models.

Parameters

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

6.3.2 Function Documentation

6.3 Network counters 27

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

6.3 Network counters 29

Definition at line 693 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()

Calculates the logit intercept for the DEFM model.

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Parameters

counters	A pointer to the DEFMCounters object.
n_y	The number of response variables.
which	A vector of indices indicating which response variables to use. If empty, all response variables are used.
covar_index	The index of the covariate to use as the intercept.
vname	The name of the variable to use as the intercept. If empty, the intercept is set to zero.
x_names	A pointer to a vector of strings containing the names of the covariates.
y_names	A pointer to a vector of strings containing the names of the response variables.

Definition at line 170 of file counters.hpp.

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.

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6.3.2.26 counter_odegree15() [2/2]

Definition at line 834 of file network.hpp.

6.3.2.27 counter_ones()

Prevalence of ones.

Parameters

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

Definition at line 89 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 289 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 611 of file counters.hpp.

6.3.2.32 counter_ttriads() [1/2]

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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.

6.3.2.35 rule_constrain_support()

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 820 of file counters.hpp.

6.3.2.36 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 761 of file counters.hpp.

6.3.2.37 rules_markov_fixed()

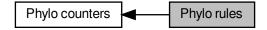
Number of edges.

Definition at line 736 of file counters.hpp.

6.4 Phylo rules

Rules for phylogenetic modeling.

Collaboration diagram for Phylo rules:



Classes

• class DEFMRuleDynData

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Functions

- DEFMData::DEFMData ()
- DEFMData::DEFMData (DEFMArray *array_, const double *covariates_, size_t obs_start_, size_t X_ncol_, size_t X_nrow_, bool column_major_)

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_, size_t pos_, size_t lb_ ← , size_t ub_)
- DEFMRuleDynData::~DEFMRuleDynData ()
- const double DEFMRuleDynData::operator() () const
- void rule_leafs (PhyloSupport *support)
- 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.

· bool DEFMData::column major

Vector indicating which covariates are included in the model.

std::vector< size_t > DEFMCounterData::indices

- std::vector< double > DEFMCounterData::numbers
- std::vector< bool > DEFMCounterData::logical
- · bool DEFMCounterData::is motif

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
- size t DEFMRuleDynData::pos
- size_t DEFMRuleDynData::lb
- size_t DEFMRuleDynData::ub

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

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 186 of file defm-types.hpp.

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6.4.2.2 DEFMCounters

typedef barry::Counters<DEFMArray, DEFMCounterData> DEFMCounters

Definition at line 187 of file defm-types.hpp.

6.4.2.3 DEFMModel

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

Definition at line 190 of file defm-types.hpp.

6.4.2.4 DEFMRule

typedef barry::Rule<DEFMArray, DEFMRuleData> DEFMRule

Definition at line 193 of file defm-types.hpp.

6.4.2.5 DEFMRuleDyn

typedef barry::Rule<DEFMArray, DEFMRuleDynData> DEFMRuleDyn

Definition at line 195 of file defm-types.hpp.

6.4.2.6 DEFMRules

typedef barry::Rules<DEFMArray, DEFMRuleData> DEFMRules

Definition at line 194 of file defm-types.hpp.

6.4.2.7 DEFMRulesDyn

typedef barry::Rules<DEFMArray, DEFMRuleDynData> DEFMRulesDyn

Definition at line 196 of file defm-types.hpp.

6.4.2.8 DEFMStatsCounter

```
typedef barry::StatsCounter<DEFMArray, DEFMCounterData> DEFMStatsCounter
```

Definition at line 189 of file defm-types.hpp.

6.4.2.9 **DEFMSupport**

```
typedef barry::Support<DEFMArray, DEFMCounterData, DEFMRuleData,DEFMRuleDynData> DEFMSupport
```

Definition at line 188 of file defm-types.hpp.

6.4.3 Function Documentation

6.4.3.1 at()

6.4.3.2 DEFMCounterData() [1/2]

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

Definition at line 74 of file defm-types.hpp.

6.4.3.3 DEFMCounterData() [2/2]

Definition at line 75 of file defm-types.hpp.

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6.4.3.4 DEFMData() [1/2]

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

Definition at line 28 of file defm-types.hpp.

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

Constructor.

Parameters

covariates⇔	Pointer to the attribute data.
_	
obs_←	Location of the current observation in the covariates vector
start_	
X_ncol_	Number of columns (covariates.)

Definition at line 37 of file defm-types.hpp.

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

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

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

6.4.3.7 DEFMRuleData() [2/3]

Definition at line 112 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 106 of file defm-types.hpp.

6.4.3.9 DEFMRuleDynData()

Definition at line 166 of file defm-types.hpp.

6.4.3.10 idx() [1/2]

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

6.4.3.11 idx() [2/2]

Definition at line 101 of file defm-types.hpp.

6.4.3.12 is_true() [1/2]

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

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6.4.3.13 is_true() [2/2]

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

6.4.3.14 ncol()

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

Definition at line 130 of file defm-types.hpp.

6.4.3.15 nrow()

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

Definition at line 134 of file defm-types.hpp.

6.4.3.16 num() [1/2]

Definition at line 84 of file defm-types.hpp.

6.4.3.17 num() [2/2]

```
double DEFMRuleData::num ( \label{eq:size_ti} \mbox{size\_t $i$ ) const [inline]}
```

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

6.4.3.18 operator()() [1/2]

```
const double DEFMRuleDynData::operator() ( ) const [inline]
```

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

6.4.3.19 operator()() [2/2]

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

Parameters

i	
j	

Returns

double

Definition at line 120 of file defm-types.hpp.

6.4.3.20 print()

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

Definition at line 138 of file defm-types.hpp.

6.4.3.21 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 2087 of file counters.hpp.

6.4 Phylo rules 45

6.4.3.22 rule_leafs()

Definition at line 2052 of file counters.hpp.

6.4.3.23 \sim DEFMCounterData()

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

Definition at line 87 of file defm-types.hpp.

6.4.3.24 \sim DEFMData()

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

Definition at line 60 of file defm-types.hpp.

6.4.3.25 ∼DEFMRuleDynData()

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

Definition at line 173 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 column_major

```
bool DEFMData::column_major
```

Vector indicating which covariates are included in the model.

Definition at line 26 of file defm-types.hpp.

6.4.4.3 counts

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

Definition at line 161 of file defm-types.hpp.

6.4.4.4 covar_sort

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

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

6.4.4.5 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.

6.4.4.6 covariates

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

Vector of covariates (complete vector)

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

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6.4.4.7 indices [1/2]

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

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

6.4.4.8 indices [2/2]

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

Definition at line 95 of file defm-types.hpp.

6.4.4.9 init

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

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

6.4.4.10 is_motif

bool DEFMCounterData::is_motif

If false, then is a logit intercept.

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

6.4.4.11 lb

```
size_t DEFMRuleDynData::lb
```

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

6.4.4.12 logical [1/2]

std::vector< bool > DEFMCounterData::logical

Definition at line 71 of file defm-types.hpp.

6.4.4.13 logical [2/2]

std::vector< bool > DEFMRuleData::logical

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

6.4.4.14 numbers [1/2]

std::vector< double > DEFMCounterData::numbers

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

6.4.4.15 numbers [2/2]

std::vector< double > DEFMRuleData::numbers

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

6.4.4.16 obs_start

size_t DEFMData::obs_start

Index of the observation in the data.

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

6.4.4.17 pos

size_t DEFMRuleDynData::pos

Definition at line 162 of file defm-types.hpp.

6.4.4.18 ub

size_t DEFMRuleDynData::ub

Definition at line 164 of file defm-types.hpp.

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6.4.4.19 X_ncol

```
size_t DEFMData::X_ncol
```

Number of columns in the array of covariates.

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

6.4.4.20 X_nrow

```
size_t DEFMData::X_nrow
```

Number of rows in the array of covariates.

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

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.

 $\bullet \ \ void\ counter_prop_genes_changing\ (PhyloCounters\ *counters,\ size_t\ duplication=Geese::etype_default)$

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)
- $\bullet \ \ void\ counter_longest\ (PhyloCounters\ *counters,\ size_t\ duplication=Geese::etype_default)$

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 counters_gains_from_0 (PhyloCounters *counters_std::vector< size_t > pfun_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.)

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>).

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6.5.2 Function Documentation

6.5.2.1 counter_co_opt()

Function co-opting.

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

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

This algorithm implements the change statistic.

Definition at line 1299 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 794 of file counters.hpp.

6.5.2.3 counter_gains()

Functional gains for a specific function (nfun).

Definition at line 99 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 1633 of file counters.hpp.

6.5.2.5 counter_gains_k_offspring()

k genes gain function nfun

Definition at line 159 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 231 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 1397 of file counters.hpp.

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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 1517 of file counters.hpp.

6.5.2.9 counter_longest()

Longest branch mutates (either by gain or by loss)

Definition at line 851 of file counters.hpp.

6.5.2.10 counter_loss()

Total count of losses for an specific function.

Definition at line 594 of file counters.hpp.

6.5.2.11 counter_maxfuns()

Cap the number of functions per gene.

Definition at line 532 of file counters.hpp.

6.5.2.12 counter_neofun()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1021 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 1166 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 646 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 61 of file counters.hpp.

6.5 Phylo counters 55

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 1699 of file counters.hpp.

6.5.2.17 counter_overall_loss()

Overall functional loss.

Definition at line 484 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 $x(a)^3(1-x(b))^3 + x(b)^3(1-x(a))^3 + x(a)^3 * x(b)^3 + (1-x(a))^3 * (1-x(b))^3$

Definition at line 1951 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 1102 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 1747 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 1812 of file counters.hpp.

6.5.2.22 counter_preserve_pseudogene()

Keeps track of how many pairs of genes preserve pseudostate.

Definition at line 300 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 382 of file counters.hpp.

6.5 Phylo counters 57

6.5.2.24 counter_subfun()

Total count of Sub-functionalization events.

It requires to specify data = {funA, funB}

Definition at line 705 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_)
- ∼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
- void print_n (size_t nrow, size_t ncol, 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_
General Bet by	לאשׁם 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)
- 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	ck_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

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

Baseline class for binary arrays.

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

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]

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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 241 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]

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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 print_n()

8.1.3.39 reserve()

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

8.1.3.40 resize()

8.1.3.41 rm_cell()

8.1.3.42 row()

8.1.3.43 set_data()

Set the data object.

Parameters

data_	
delete_⊸	
data_	

8.1.3.44 swap_cells()

8.1.3.45 swap_cols()

8.1.3.46 swap_rows()

8.1.3.47 toggle_cell()

8.1.3.48 toggle_lock()

8.1.3.49 transpose()

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

8.1.3.50 zero_col()

8.1.3.51 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
```

8.2.1 Detailed Description

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

• bool operator== (const Cell_Type &val) const

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 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
- 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_t i) 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.

- BArrayDense (BArrayDense < Cell_Type, Data_Type > &&x) noexcept
- 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)
- $\bullet \ \, \mathsf{BArrayDense} < \mathsf{Cell_Type}, \, \mathsf{Data_Type} > \& \, \mathsf{operator}\text{--=} \, (\mathsf{const} \, \, \mathsf{std} \text{::pair} < \, \mathsf{size_t}, \, \mathsf{size_t} > \& \mathsf{coords})$
- 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)
- void insert cell (size ti, size tj, Cell Type v, bool check bounds, bool)

- 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

- 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.

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

8.4.2.4 BArrayDense() [4/6]

Edgelist with no data (simpler)

Definition at line 86 of file barraydense-meat.hpp.

8.4.2.5 BArrayDense() [5/6]

Copy constructor.

Definition at line 135 of file barraydense-meat.hpp.

8.4.2.6 BArrayDense() [6/6]

Move operator.

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

8.4.2.7 ~BArrayDense()

```
template<typename Cell_Type , typename Data_Type >
BArrayDense< Cell_Type, Data_Type >::~BArrayDense [inline]
```

Definition at line 310 of file barraydense-meat.hpp.

8.4.3 Member Function Documentation

8.4.3.1 clear()

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

Definition at line 911 of file barraydense-meat.hpp.

8.4.3.2 col() [1/2]

Definition at line 504 of file barraydense-meat.hpp.

8.4.3.3 col() [2/2]

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

8.4.3.4 colsum()

Definition at line 1014 of file barraydense-meat.hpp.

8.4.3.5 D() [1/2]

```
template<typename Cell_Type , typename Data_Type >
Data_Type & BArrayDense< Cell_Type, Data_Type >::D [inline]
```

Definition at line 345 of file barraydense-meat.hpp.

8.4.3.6 D() [2/2]

```
template<typename Cell_Type , typename Data_Type >
const Data_Type & BArrayDense< Cell_Type, Data_Type >::D [inline]
```

Definition at line 350 of file barraydense-meat.hpp.

8.4.3.7 D_ptr() [1/2]

```
template<typename Cell_Type , typename Data_Type >
Data_Type * BArrayDense< Cell_Type, Data_Type >::D_ptr [inline]
```

Definition at line 335 of file barraydense-meat.hpp.

8.4.3.8 **D_ptr()** [2/2]

```
template<typename Cell_Type , typename Data_Type >
const Data_Type * BArrayDense< Cell_Type, Data_Type >::D_ptr [inline]
```

Definition at line 340 of file barraydense-meat.hpp.

8.4.3.9 default_val()

```
template<typename Cell_Type , typename Data_Type >
Cell< Cell_Type > BArrayDense< Cell_Type, Data_Type >::default_val [inline]
```

Definition at line 577 of file barraydense-meat.hpp.

8.4.3.10 get_cell()

Definition at line 376 of file barraydense-meat.hpp.

8.4.3.11 get_col_vec() [1/2]

Definition at line 424 of file barraydense-meat.hpp.

8.4.3.12 get_col_vec() [2/2]

Definition at line 442 of file barraydense-meat.hpp.

8.4.3.13 get_data()

```
template<typename Cell_Type , typename Data_Type >
const std::vector< Cell_Type > & BArrayDense< Cell_Type, Data_Type >::get_data [inline]
```

Definition at line 1004 of file barraydense-meat.hpp.

8.4.3.14 get_entries()

```
template<typename Cell_Type , typename Data_Type >
Entries< Cell_Type > BArrayDense< Cell_Type, Data_Type >::get_entries [inline]
```

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>
```

Definition at line 516 of file barraydense-meat.hpp.

8.4.3.15 get_row_vec() [1/2]

Definition at line 391 of file barraydense-meat.hpp.

8.4.3.16 get_row_vec() [2/2]

Definition at line 409 of file barraydense-meat.hpp.

8.4.3.17 insert cell() [1/2]

Definition at line 703 of file barraydense-meat.hpp.

8.4.3.18 insert_cell() [2/2]

Definition at line 669 of file barraydense-meat.hpp.

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()

Definition at line 545 of file barraydense-meat.hpp.

8.4.3.21 ncol()

```
template<typename Cell_Type , typename Data_Type >
size_t BArrayDense< Cell_Type, Data_Type >::ncol [inline], [noexcept]
```

Definition at line 562 of file barraydense-meat.hpp.

8.4.3.22 nnozero()

```
template<typename Cell_Type , typename Data_Type >
size_t BArrayDense< Cell_Type, Data_Type >::nnozero [inline], [noexcept]
```

Definition at line 566 of file barraydense-meat.hpp.

8.4.3.23 nrow()

```
template<typename Cell_Type , typename Data_Type >
size_t BArrayDense< Cell_Type, Data_Type >::nrow [inline], [noexcept]
```

Definition at line 558 of file barraydense-meat.hpp.

8.4.3.24 operator()() [1/2]

Definition at line 621 of file barraydense-meat.hpp.

8.4.3.25 operator()() [2/2]

Definition at line 632 of file barraydense-meat.hpp.

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]

Definition at line 582 of file barraydense-meat.hpp.

8.4.3.30 operator-=() [1/3]

8.4.3.31 operator-=() [2/3]

8.4.3.32 operator-=() [3/3]

Definition at line 601 of file barraydense-meat.hpp.

8.4.3.33 operator/=()

8.4.3.34 operator=() [1/2]

Move assignment.

Definition at line 247 of file barraydense-meat.hpp.

8.4.3.35 operator=() [2/2]

Assignment constructor.

Definition at line 179 of file barraydense-meat.hpp.

8.4.3.36 operator==()

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

8.4.3.37 out_of_range()

Definition at line 355 of file barraydense-meat.hpp.

8.4.3.38 print()

Definition at line 970 of file barraydense-meat.hpp.

8.4.3.39 reserve()

```
template<typename Cell_Type , typename Data_Type >
void BArrayDense< Cell_Type, Data_Type >::reserve [inline]
```

Definition at line 961 of file barraydense-meat.hpp.

8.4.3.40 resize()

Definition at line 925 of file barraydense-meat.hpp.

8.4.3.41 rm_cell()

Definition at line 646 of file barraydense-meat.hpp.

8.4.3.42 row() [1/2]

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

8.4.3.43 row() [2/2]

Definition at line 463 of file barraydense-meat.hpp.

8.4.3.44 rowsum()

Definition at line 1009 of file barraydense-meat.hpp.

8.4.3.45 set_data()

Set the data object.

Parameters

data_	
delete_⊸	
data_	

Definition at line 319 of file barraydense-meat.hpp.

8.4.3.46 swap_cells()

```
template<typename Cell_Type , typename Data_Type >
void BArrayDense< Cell_Type, Data_Type >::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 ) [inline]
```

Definition at line 734 of file barraydense-meat.hpp.

8.4.3.47 swap_cols()

Definition at line 818 of file barraydense-meat.hpp.

8.4.3.48 swap_rows()

Definition at line 790 of file barraydense-meat.hpp.

8.4.3.49 toggle_cell()

Definition at line 771 of file barraydense-meat.hpp.

8.4.3.50 toggle_lock()

8.4.3.51 transpose()

```
template<typename Cell_Type , typename Data_Type >
void BArrayDense< Cell_Type, Data_Type >::transpose [inline]
```

Definition at line 883 of file barraydense-meat.hpp.

8.4.3.52 zero_col()

Definition at line 864 of file barraydense-meat.hpp.

8.4.3.53 zero_row()

Definition at line 845 of file barraydense-meat.hpp.

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

- BArrayDenseCell (BArrayDense< Cell_Type, Data_Type > *Array_, size_t i_, size_t j_, bool check_←
 bounds=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 112 of file barraydensecell-meat.hpp.

8.5.3.2 operator*=()

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

8.5.3.3 operator+=()

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

8.5.3.4 operator-=()

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

8.5.3.5 operator/=()

Definition at line 95 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 31 of file barraydensecell-meat.hpp.

8.5.3.8 operator==()

Definition at line 117 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
- class BArrayDenseCell_const< Cell_Type, 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 \sim 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.

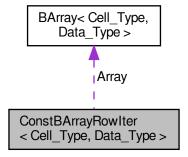
The documentation for this class was generated from the following file:

• 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:

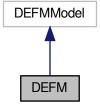
• include/barry/counters-bones.hpp

8.20 DEFM Class Reference 143

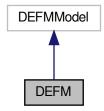
8.20 DEFM Class Reference

#include <defm-bones.hpp>

Inheritance diagram for DEFM:



Collaboration diagram for DEFM:



Public Member Functions

- DEFM (int *id, int *y, double *x, size_t id_length, size_t y_ncol, size_t x_ncol, size_t m_order, bool copy_
 data=true, bool column_major=true)
- DEFMModel & get_model ()
- void init ()
- 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 ()
bool get_column_major () const noexcept
```

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()

```
DEFM::DEFM (
    int * id,
    int * y,
    double * x,
    size_t id_length,
    size_t y_ncol,
    size_t x_ncol,
    size_t m_order,
    bool copy_data = true,
    bool column_major = true ) [inline]
```

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

8.20.3 Member Function Documentation

```
8.20.3.1 get_column_major()
```

```
bool DEFM::get_column_major ( ) const [inline], [noexcept]
```

Definition at line 458 of file defm-meat.hpp.

8.20.3.2 get_ID()

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

Definition at line 316 of file defm-meat.hpp.

8.20 DEFM Class Reference 145

8.20.3.3 get_m_order()

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

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

8.20.3.4 get_model()

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

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

8.20.3.5 get_n_covars()

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

Definition at line 296 of file defm-meat.hpp.

8.20.3.6 get_n_obs()

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

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

8.20.3.7 get_n_rows()

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

Definition at line 306 of file defm-meat.hpp.

8.20.3.8 get_n_y()

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

Definition at line 286 of file defm-meat.hpp.

8.20.3.9 get_X()

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

Definition at line 321 of file defm-meat.hpp.

8.20.3.10 get_X_names()

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

Definition at line 431 of file defm-meat.hpp.

8.20.3.11 get_Y()

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

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

8.20.3.12 get_Y_names()

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

Definition at line 427 of file defm-meat.hpp.

8.20.3.13 init()

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

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

8.20.3.14 is_motif()

```
std::vector< bool > DEFM::is_motif ( ) [inline]
```

Definition at line 448 of file defm-meat.hpp.

8.20 DEFM Class Reference 147

8.20.3.15 logodds()

Definition at line 365 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 327 of file defm-meat.hpp.

8.20.3.17 print()

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

Definition at line 435 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 410 of file defm-meat.hpp.

8.20.3.19 simulate()

```
void DEFM::simulate (
          std::vector< double > par,
          int * y_out ) [inline]
```

Definition at line 39 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_t i) 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 66 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 ()
- DEFMData (DEFMArray *array_, const double *covariates_, size_t obs_start_, size_t X_ncol_, size_t X_
 nrow_, bool column_major_)

Constructor.

• double operator() (size_t i, size_t j) const

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

- double at (size ti, size ti) 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 tobs 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.

· bool column_major

Vector indicating which covariates are included in the model.

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>
```

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 91 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>
```

Public Member Functions

- DEFMRuleDynData (const std::vector< double > *counts_, size_t pos_, size_t lb_, size_t ub_)
- ~DEFMRuleDynData ()
- const double operator() () const

Public Attributes

- const std::vector< double > * counts
- size_t pos
- size_t lb
- size_t ub

8.24.1 Detailed Description

Definition at line 159 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

```
template<typename Cell_Type> class Entries< Cell_Type>
```

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 ∼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 153

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< 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
- size_t parse_polytomies (bool verb=true, std::vector< size_t > *dist=nullptr) const noexcept

Check polytomies and return the largest.

· void print () const

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.

8.26 Flock Class Reference 155

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 246 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< double > * Flock::get_stats_support ( ) [inline]
```

Definition at line 117 of file flock-meat.hpp.

8.26.3.6 get_stats_target()

```
std::vector< std::vector< 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	When true (default) will compute the likelihood using the 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 189 of file flock-meat.hpp.

8.26 Flock Class Reference 157

8.26.3.11 nleafs()

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

Definition at line 217 of file flock-meat.hpp.

8.26.3.12 nnodes()

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

Definition at line 203 of file flock-meat.hpp.

8.26.3.13 nterms()

```
size_t Flock::nterms ( ) const [inline]
```

Definition at line 231 of file flock-meat.hpp.

8.26.3.14 ntrees()

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

Definition at line 196 of file flock-meat.hpp.

8.26.3.15 operator()()

Access the i-th geese element.

Parameters

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

Returns

Geese*

Definition at line 324 of file flock-meat.hpp.

8.26.3.16 parse_polytomies()

Check polytomies and return the largest.

Definition at line 253 of file flock-meat.hpp.

8.26.3.17 print()

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

Definition at line 280 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 239 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 \sim 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 163

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, size_t ncores=1u, bool no_update_pset_probs=false)
- 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 print_nodes () const
- 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.

std::vector< size_t > get_annotations () const

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

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- 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_state_id
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
```

8.28.1 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_\leftrightarrow 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 114 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 230 of file geese-meat-constructors.hpp.

8.28.2.4 Geese() [4/4]

Definition at line 309 of file geese-meat-constructors.hpp.

8.28.2.5 ∼Geese()

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

Definition at line 119 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 361 of file geese-meat.hpp.

8.28.3.2 calc_sequence()

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

Definition at line 317 of file geese-meat.hpp.

8.28.3.3 colnames()

```
std::vector< std::string > Geese::colnames ( ) const [inline]
```

Names of the terms in the model.

Definition at line 482 of file geese-meat.hpp.

8.28.3.4 get_annotated_nodes()

```
\verb|std::vector| < \verb|size_t| > \verb|Geese::get_annotated_nodes| ( ) | const | [inline]|
```

Returns the ids of the nodes with at least one annotation.

Definition at line 771 of file geese-meat.hpp.

8.28.3.5 get_annotations()

```
\verb|std::vector| < \verb|size_t| > \verb|Geese::get_annotations| ( ) | const | [inline]|
```

Returns the annotations of the nodes with at least one annotation.

Definition at line 794 of file geese-meat.hpp.

8.28.3.6 get_counters()

```
PhyloCounters * Geese::get_counters ( ) [inline]
```

Definition at line 754 of file geese-meat.hpp.

8.28.3.7 get_model()

```
PhyloModel * Geese::get_model ( ) [inline]
```

Definition at line 759 of file geese-meat.hpp.

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8.28.3.8 get_probabilities()

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

Definition at line 409 of file geese-meat.hpp.

8.28.3.9 get_rengine()

```
std::mt19937 * Geese::get_rengine ( ) [inline]
```

Definition at line 749 of file geese-meat.hpp.

8.28.3.10 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^{^{\land}}P.
```

Definition at line 767 of file geese-meat.hpp.

8.28.3.11 get_support_fun()

```
PhyloSupport * Geese::get_support_fun ( ) [inline]
```

Definition at line 763 of file geese-meat.hpp.

8.28.3.12 inherit_support()

Definition at line 260 of file geese-meat.hpp.

8.28.3.13 init()

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

8.28.3.14 init_node()

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

8.28.3.15 likelihood()

Definition at line 103 of file geese-meat-likelihood.hpp.

8.28.3.16 likelihood_exhaust()

```
double Geese::likelihood_exhaust ( {\tt const\ std::vector<\ double\ >\ \&\ par\ )} \quad [{\tt inline}]
```

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

8.28.3.17 nannotations()

```
std::vector< size_t > Geese::nannotations ( ) const [inline], [noexcept]
```

Number of annotations.

Definition at line 473 of file geese-meat.hpp.

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8.28.3.18 nfuns()

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

Number of functions analyzed.

Definition at line 430 of file geese-meat.hpp.

8.28.3.19 nleafs()

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

Number of leaf.

Definition at line 444 of file geese-meat.hpp.

8.28.3.20 nnodes()

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

Number of nodes (interior + leaf)

Definition at line 437 of file geese-meat.hpp.

8.28.3.21 nterms()

```
size_t Geese::nterms ( ) const [inline]
```

Number of terms included.

Definition at line 456 of file geese-meat.hpp.

8.28.3.22 observed_counts()

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

Definition at line 524 of file geese-meat.hpp.

8.28.3.23 operator=() [1/2]

8.28.3.24 operator=() [2/2]

8.28.3.25 parse_polytomies()

Check polytomies and return the largest.

Definition at line 489 of file geese-meat.hpp.

8.28.3.26 predict()

Definition at line 279 of file geese-meat-predict.hpp.

8.28.3.27 predict_backend()

< True if the array belongs to the set

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

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8.28.3.28 predict_exhaust()

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

Definition at line 5 of file geese-meat-predict_exhaust.hpp.

8.28.3.29 predict_exhaust_backend()

Definition at line 47 of file geese-meat-predict_exhaust.hpp.

8.28.3.30 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.

8.28.3.31 print()

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

Prints information about the GEESE.

Definition at line 658 of file geese-meat.hpp.

8.28.3.32 print_nodes()

```
void Geese::print_nodes ( ) const [inline]
```

Definition at line 676 of file geese-meat.hpp.

8.28.3.33 print_observed_counts()

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

Definition at line 595 of file geese-meat.hpp.

8.28.3.34 set_seed()

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

8.28.3.35 simulate()

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

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

8.28.3.36 support_size()

```
size_t Geese::support_size ( ) const [inline], [noexcept]
```

Number of unique sets of sufficient stats.

Definition at line 463 of file geese-meat.hpp.

8.28.3.37 update_annotations()

Definition at line 288 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 153 of file geese-bones.hpp.

8.28.4.2 delete_support

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

Definition at line 154 of file geese-bones.hpp.

8.28.4.3 etype_default

```
const size_t Geese::etype_default = 1ul [static]
```

Definition at line 167 of file geese-bones.hpp.

8.28.4.4 etype_duplication

```
const size_t Geese::etype_duplication = 1ul [static]
```

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

8.28.4.5 etype_either

```
const size_t Geese::etype_either = 2ul [static]
```

Definition at line 170 of file geese-bones.hpp.

8.28.4.6 etype_speciation

```
const size_t Geese::etype_speciation = Oul [static]
```

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

8.28.4.7 initialized

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

Definition at line 152 of file geese-bones.hpp.

8.28.4.8 map_to_state_id

```
barry::MapVec_type< size_t > Geese::map_to_state_id
```

Definition at line 144 of file geese-bones.hpp.

8.28.4.9 nfunctions

```
size_t Geese::nfunctions
```

Definition at line 141 of file geese-bones.hpp.

8.28.4.10 nodes

```
std::map< size_t, Node > Geese::nodes
```

Definition at line 142 of file geese-bones.hpp.

8.28.4.11 pset_loc

```
std::vector< std::vector< size_t > > > Geese::pset_loc
```

Locations of columns.

Definition at line 145 of file geese-bones.hpp.

8.28.4.12 reduced_sequence

```
std::vector< size_t > Geese::reduced_sequence
```

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

8.28.4.13 sequence

std::vector< size_t > Geese::sequence

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

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

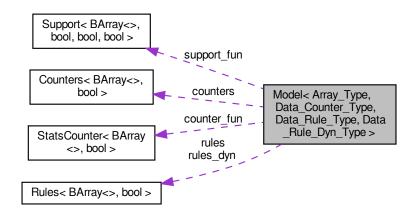
- include/barry/models/geese/geese-bones.hpp
- include/barry/models/geese/geese-meat-constructors.hpp
- include/barry/models/geese/geese-meat-likelihood.hpp
- include/barry/models/geese/geese-meat-likelihood_exhaust.hpp
- include/barry/models/geese/geese-meat-predict.hpp
- include/barry/models/geese/geese-meat-predict_exhaust.hpp
- 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 update_normalizing_constants (const std::vector< double > ¶ms, BARRY_NCORES_ARG(=1), int i=-1)

Computes the normalizing constant for a given set of parameters.

- void update_likelihoods (const std::vector< double > ¶ms,)
- void update pset probs (const std::vector< double > ¶ms, BARRY NCORES ARG(=1), int i=-1)
- void set rengine (std::mt19937 *rengine , bool delete =false)
- void set seed (size ts)
- 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 > ¶ms={})
- Array_Type sample (const size_t &i, const std::vector< double > ¶ms)
- double conditional_prob (const Array_Type &Array_, const std::vector< double > ¶ms, 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.

- $\bullet \ \ \mathsf{void} \ \ \mathsf{add_rule} \ (\mathsf{Rule} {<} \ \mathsf{Array_Type}, \ \mathsf{Data_Rule_Type} > \& \mathsf{rule}) \\$
- 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 add_rule_dyn (Rule_fun_type< Array_Type, Data_Rule_Dyn_Type > count_fun_, Data_Rule_Dyn
 Type data)
- 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 > ¶ms, const size_t &i, bool as_log=false, bool no_

 update normconst=false)
- double likelihood (const std::vector< double > ¶ms, const Array_Type &Array_, int i=-1, bool as_←
 log=false, bool no update normconst=false)
- double likelihood (const std::vector< double > ¶ms, const std::vector< double > &target_, const size t &i, bool as log=false, bool no update normconst=false)
- double likelihood (const std::vector < double > ¶ms, const double *target_, const size_t &i, bool as
 — log=false, bool no_update_normconst=false)
- double likelihood_total (const std::vector < double > ¶ms, bool as_log=false, BARRY_NCORES_ARG(=2), bool no_update_normconst=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.

- const std::vector< double > & get normalizing constants () const
- const std::vector< double > & get_likelihoods () const
- const std::vector< Array_Type > * get_pset (const size_t &i)
- const 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< double > * get_stats_support ()

Sufficient statistics of the support(s)

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< double > stats support

Sufficient statistics of the model (support)

std::vector < size_t > stats_support_sizes

Number of vectors included in the support.

std::vector< size_t > stats_support_sizes_acc

Accumulated number of vectors included in the 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< double > stats likelihood
- 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< double > pset_stats

Statistics of the support(s)

std::vector< double > pset_probs

Probabilities of the support(s)

• std::vector< size_t > pset_sizes

Number of vectors included in the support.

std::vector< size_t > pset_locations

Accumulated number of vectors included in the support.

Functions to compute statistics

Arguments are recycled to save memory and computation.

- Counters < Array_Type, Data_Counter_Type > * counters
- Rules < Array_Type, Data_Rule_Type > * rules
- 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 309 of file model-meat.hpp.

8.29.2.2 Model() [2/3]

Definition at line 347 of file model-meat.hpp.

8.29.2.3 Model() [3/3]

Definition at line 389 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 187 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.

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

Definition at line 633 of file model-meat.hpp.

8.29.3.2 add_counter() [1/2]

Definition at line 508 of file model-meat.hpp.

8.29.3.3 add_counter() [2/2]

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

8.29.3.4 add_hasher()

Definition at line 552 of file model-meat.hpp.

8.29.3.5 add_rule() [1/2]

Definition at line 563 of file model-meat.hpp.

8.29.3.6 add_rule() [2/2]

8.29.3.7 add_rule_dyn() [1/2]

Definition at line 591 of file model-meat.hpp.

8.29.3.8 add_rule_dyn() [2/2]

Definition at line 600 of file model-meat.hpp.

8.29.3.9 colnames()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< std::string > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Counter_Type >::colnames [inline]
```

Definition at line 1389 of file model-meat.hpp.

8.29.3.10 conditional_prob()

Conditional probability ("Gibbs sampler")

Computes the conditional probability of observing $P\{Y(i,j) = | Y^{\land}C, \text{ theta}\}$, 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

Returns

double The conditional probability

Definition at line 1689 of file model-meat.hpp.

8.29.3.11 gen_key()

Definition at line 501 of file model-meat.hpp.

8.29.3.12 get_arrays2support()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
std::vector< size_t > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn
_Type >::qet_arrays2support [inline]
```

Definition at line 1778 of file model-meat.hpp.

8.29.3.13 get_counters()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
Counters< Array_Type, Data_Counter_Type > * Model< Array_Type, Data_Counter_Type, Data_Rule
_Type, Data_Rule_Dyn_Type >::get_counters [inline]
```

Definition at line 1727 of file model-meat.hpp.

8.29.3.14 get_likelihoods()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
const std::vector< double > & Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_←
Rule_Dyn_Type >::get_likelihoods [inline]
```

Definition at line 1178 of file model-meat.hpp.

8.29.3.15 get_normalizing_constants()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
const std::vector< double > & Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_←
Rule_Dyn_Type >::get_normalizing_constants [inline]
```

Definition at line 1165 of file model-meat.hpp.

8.29.3.16 get_pset()

Definition at line 1186 of file model-meat.hpp.

8.29.3.17 get_pset_arrays()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< std::vector< Array_Type > > * Model< Array_Type, Data_Counter_Type, Data_Rule_\Lorenty
Type, Data_Rule_Dyn_Type >::get_pset_arrays [inline]
```

Definition at line 1785 of file model-meat.hpp.

8.29.3.18 get_pset_locations()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< size_t > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn 
_Type >::get_pset_locations [inline]
```

Definition at line 1823 of file model-meat.hpp.

8.29.3.19 get_pset_probs()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< double > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn
_Type >::get_pset_probs [inline]
```

Definition at line 1802 of file model-meat.hpp.

8.29.3.20 get_pset_sizes()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< size_t > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn 
_Type >::get_pset_sizes [inline]
```

Definition at line 1812 of file model-meat.hpp.

8.29.3.21 get_pset_stats() [1/2]

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< double > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn 
_Type >::get_pset_stats [inline]
```

Statistics of the support(s)

Definition at line 1791 of file model-meat.hpp.

8.29.3.22 get_pset_stats() [2/2]

Definition at line 1200 of file model-meat.hpp.

8.29.3.23 get_rengine()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
const std::mt19937 * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type
>::get_rengine [inline]
```

Definition at line 1722 of file model-meat.hpp.

8.29.3.24 get_rules()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
Rules< Array_Type, Data_Rule_Type > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_rules [inline]
```

Definition at line 1732 of file model-meat.hpp.

8.29.3.25 get_rules_dyn()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
Rules< Array_Type, Data_Rule_Dyn_Type > * Model< Array_Type, Data_Counter_Type, Data_Rule_\Log_Type, Data_Rule_\Log_Type, Data_Rule_\Log_Type >::get_rules_dyn [inline]
```

Definition at line 1737 of file model-meat.hpp.

8.29.3.26 get_stats_support()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< double > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn 
_Type >::get_stats_support [inline]
```

Sufficient statistics of the support(s)

Definition at line 1755 of file model-meat.hpp.

8.29.3.27 get_stats_support_sizes()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< size_t > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn 
_Type >::get_stats_support_sizes [inline]
```

Number of vectors included in the support.

Definition at line 1763 of file model-meat.hpp.

8.29.3.28 get_stats_support_sizes_acc()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< size_t > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn
_Type >::get_stats_support_sizes_acc [inline]
```

Accumulated number of vectors included in the support.

Definition at line 1771 of file model-meat.hpp.

8.29.3.29 get_stats_target()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
std::vector< std::vector< double > > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_stats_target [inline]
```

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.

Definition at line 1748 of file model-meat.hpp.

8.29.3.30 get_support_fun()

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

Definition at line 1743 of file model-meat.hpp.

8.29.3.31 likelihood() [1/4]

Definition at line 857 of file model-meat.hpp.

8.29.3.32 likelihood() [2/4]

Definition at line 1007 of file model-meat.hpp.

8.29.3.33 likelihood() [3/4]

Definition at line 810 of file model-meat.hpp.

8.29.3.34 likelihood() [4/4]

Definition at line 941 of file model-meat.hpp.

8.29.3.35 likelihood_total()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarray = bool, typename Data_\leftarray = bool, typename Data_Rule_Dyn_Type = bool>
double Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::likelihood\leftarray_total (

const std::vector< double > & params,
bool as_log = false,
BARRY_NCORES_ARG(=2) ,
bool no_update_normconst = false ) [inline]
```

Definition at line 1079 of file model-meat.hpp.

8.29.3.36 nrules()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
size_t Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::nrules
[inline], [noexcept]
```

Definition at line 1359 of file model-meat.hpp.

8.29.3.37 nrules_dyn()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
size_t Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::nrules_dyn
[inline], [noexcept]
```

Definition at line 1367 of file model-meat.hpp.

8.29.3.38 nterms()

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

Definition at line 1348 of file model-meat.hpp.

8.29.3.39 operator=()

Definition at line 439 of file model-meat.hpp.

8.29.3.40 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 1252 of file model-meat.hpp.

8.29.3.41 print stats()

Definition at line 1212 of file model-meat.hpp.

8.29.3.42 sample() [1/2]

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

Definition at line 1476 of file model-meat.hpp.

8.29.3.43 sample() [2/2]

Definition at line 1406 of file model-meat.hpp.

8.29.3.44 set counters()

Definition at line 534 of file model-meat.hpp.

8.29.3.45 set_rengine()

```
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_rengine (
    std::mt19937 * rengine_,
    bool delete_ = false ) [inline]
```

Definition at line 157 of file model-bones.hpp.

8.29.3.46 set_rules()

Definition at line 573 of file model-meat.hpp.

8.29.3.47 set_rules_dyn()

Definition at line 615 of file model-meat.hpp.

8.29.3.48 set_seed()

Definition at line 167 of file model-bones.hpp.

8.29.3.49 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	

Definition at line 1835 of file model-meat.hpp.

8.29.3.50 size()

 $\label{template} $$ $$ template < typename \ Data_Counter_Type \ , \ typename \ Data_Rule_Type \ , \ typename \ Data_Rule_Dyn_Type \ > $$$

```
size_t Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::size [inline],
[noexcept]
```

Definition at line 1331 of file model-meat.hpp.

8.29.3.51 size_unique()

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

Definition at line 1339 of file model-meat.hpp.

8.29.3.52 store_psets()

```
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 >::store_psets
[inline], [noexcept]
```

Definition at line 495 of file model-meat.hpp.

8.29.3.53 support_size()

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

Definition at line 1375 of file model-meat.hpp.

8.29.3.54 transform_model()

8.29.3.55 update_likelihoods()

Definition at line 186 of file model-meat.hpp.

8.29.3.56 update_normalizing_constants()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\hookleftarrow Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::update_\hookleftarrow normalizing_constants ( const std::vector< double > & params, BARRY_NCORES_ARG(=1) , int i = -1 ) [inline]
```

Computes the normalizing constant for a given set of parameters.

This function will compute the normalizing constant for a given set of parameters. It will also update the normalizing_constants member variable.

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

8.29.3.57 update_pset_probs()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarrow Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::update_pset \leftarrow _probs ( const std::vector< double > & params, BARRY_NCORES_ARG(=1) , int i = -1) [inline]
```

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

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 68 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_←
Type, Data_Rule_Dyn_Type >::counter_fun [protected]
```

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

8.29.4.3 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 >::counters [protected]
```

Definition at line 96 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 108 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 109 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 110 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 106 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 75 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 105 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 104 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 84 of file model-bones.hpp.

8.29.4.13 pset locations

```
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 >::pset_locations [protected]
```

Accumulated number of vectors included in the support.

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

8.29.4.14 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< 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 86 of file model-bones.hpp.

8.29.4.15 pset sizes

```
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 >::pset_sizes [protected]
```

Number of vectors included in the support.

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

8.29.4.16 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< 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 85 of file model-bones.hpp.

8.29.4.17 rengine

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarray_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::mt19937* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >\leftarray_regine = nullptr [protected]
```

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

8.29.4.18 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 97 of file model-bones.hpp.

8.29.4.19 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 98 of file model-bones.hpp.

8.29.4.20 stats likelihood

```
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 >::stats_likelihood [protected]
```

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

8.29.4.21 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< 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.

8.29.4.22 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 65 of file model-bones.hpp.

8.29.4.23 stats_support_sizes

```
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_sizes [protected]
```

Number of vectors included in the support.

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

8.29.4.24 stats_support_sizes_acc

```
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_sizes_acc [protected]
```

Accumulated number of vectors included in the support.

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

8.29.4.25 stats_target

```
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 >::stats_target [protected]
```

Target statistics of the model.

Definition at line 66 of file model-bones.hpp.

8.29.4.26 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 99 of file model-bones.hpp.

8.29.4.27 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 128 of file model-bones.hpp.

8.29.4.28 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 130 of file model-bones.hpp.

8.29.4.29 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 83 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 ()
- $\bullet \ \ {\tt NetCounterData} \ ({\tt const} \ {\tt std} :: {\tt vector} < {\tt size_t} > {\tt \&indices_, const} \ {\tt std} :: {\tt vector} < {\tt double} > {\tt \&numbers_)}$
- ∼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.

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

Constructor using multiple attributes.

∼NetworkData ()

Public Attributes

- bool directed = true
- std::vector< std::vector< double >> vertex attr

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]

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

Constructor using multiple attributes.

Parameters

vertex_← attr_	Vector of double vectors. The size equals to the number of attributes to be created. Each 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.

8.32 Node Class Reference 209

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)

Public Attributes

```
size_t id
```

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.

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 37 of file geese-node-bones.hpp.

8.32 Node Class Reference 211

8.32.2.2 Node() [2/5]

Definition at line 57 of file geese-node-bones.hpp.

8.32.2.3 Node() [3/5]

Definition at line 63 of file geese-node-bones.hpp.

8.32.2.4 Node() [4/5]

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

8.32.2.5 Node() [5/5]

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

8.32.2.6 ∼Node()

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

Definition at line 48 of file geese-node-bones.hpp.

8.32.3 Member Function Documentation

8.32.3.1 get_parent()

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

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

8.32.3.2 is_leaf()

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

Definition at line 112 of file geese-node-bones.hpp.

8.32.3.3 noffspring()

```
size_t Node::noffspring ( ) const [inline], [noexcept]
```

Definition at line 106 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 Node Class Reference 213

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 duplication

```
bool Node::duplication
```

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

8.32.4.5 id

```
size_t Node::id
```

Id of the node (as specified in the input)

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

8.32.4.6 narray

```
std::vector< size_t > Node::narray = {}
```

ID of the array in the model.

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

8.32.4.7 offspring

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

Offspring nodes.

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

8.32.4.8 ord

```
size_t Node::ord
```

Order in which the node was created.

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

8.32.4.9 parent

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

Parent node.

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

8.32.4.10 probability

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

The probability of observing each state.

Definition at line 29 of file geese-node-bones.hpp.

8.32.4.11 subtree_prob

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

Induced subtree probabilities.

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

8.32.4.12 visited

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

Definition at line 26 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, bool has_leaf_=false)

Public Attributes

```
    std::vector< double > blengths = {}
    std::vector< bool > states = {}
    bool duplication = true
        Whether the node is a duplication.
    bool has_leaf = false
        Whether the node has a leaf as offspring.
```

8.33.1 Detailed Description

Data definition for the PhyloArray class.

This holds basic information about a given node.

Definition at line 15 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 21 of file geese-types.hpp.

8.33.3.2 duplication

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

Whether the node is a duplication.

Definition at line 28 of file geese-types.hpp.

8.33.3.3 has_leaf

```
bool NodeData::has_leaf = false
```

Whether the node has a leaf as offspring.

Definition at line 29 of file geese-types.hpp.

8.33.3.4 states

```
std::vector< bool > NodeData::states = {}
```

State of the parent node.

Definition at line 26 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 44 of file geese-types.hpp.

8.34.2 Constructor & Destructor Documentation

8.34.2.1 PhyloCounterData() [1/2]

Definition at line 50 of file geese-types.hpp.

8.34.2.2 PhyloCounterData() [2/2]

```
PhyloCounterData::PhyloCounterData ( ) [inline]
```

Definition at line 55 of file geese-types.hpp.

8.34.3 Member Function Documentation

8.34.3.1 at()

Definition at line 57 of file geese-types.hpp.

8.34.3.2 begin()

```
std::vector< size_t >::iterator PhyloCounterData::begin ( ) [inline]
```

Definition at line 65 of file geese-types.hpp.

8.34.3.3 empty()

```
bool PhyloCounterData::empty ( ) [inline]
```

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

8.34.3.4 end()

```
std::vector< size_t >::iterator PhyloCounterData::end ( ) [inline]
```

Definition at line 66 of file geese-types.hpp.

8.34.3.5 get_counters()

```
\verb|std::vector<| double >* PhyloCounterData::get_counters () [inline]| \\
```

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

8.34.3.6 operator()()

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

8.34.3.7 operator[]()

Definition at line 59 of file geese-types.hpp.

8.34.3.8 push back()

Definition at line 61 of file geese-types.hpp.

8.34.3.9 reserve()

Definition at line 60 of file geese-types.hpp.

8.34.3.10 shrink_to_fit()

```
void PhyloCounterData::shrink_to_fit ( ) [inline]
```

Definition at line 62 of file geese-types.hpp.

8.34.3.11 size()

```
size_t PhyloCounterData::size ( ) [inline]
```

Definition at line 63 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 73 of file geese-types.hpp.

8.35.2 Constructor & Destructor Documentation

8.35.2.1 PhyloRuleDynData()

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

8.35.2.2 ~PhyloRuleDynData()

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

Definition at line 95 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 90 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 75 of file geese-types.hpp.

8.35.4.2 duplication

```
size_t PhyloRuleDynData::duplication
```

Definition at line 79 of file geese-types.hpp.

8.35.4.3 lb

```
size_t PhyloRuleDynData::lb
```

Definition at line 77 of file geese-types.hpp.

8.35.4.4 pos

```
size_t PhyloRuleDynData::pos
```

Definition at line 76 of file geese-types.hpp.

8.35.4.5 ub

```
size_t PhyloRuleDynData::ub
```

Definition at line 78 of file geese-types.hpp.

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

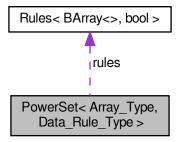
• include/barry/models/geese/geese-types.hpp

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
- $\bullet \;\; std::vector < Array_Type > ::iterator \; \frac{begin}{} \; ()$
- 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 180 of file powerset-meat.hpp.

8.36.3.2 add_rule() [2/2]

Definition at line 189 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 151 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 167 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_)
- \sim 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()

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

- Rule ()
- Rule (Rule_fun_type< Array_Type, Data_Type > fun_, Data_Type dat_, 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 180 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

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

Count stats for a single Array.

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

Definition at line 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 ~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.

- const std::vector< double > & get counts () const
- std::vector< double > * get current stats ()

List current statistics.

- · void print () const
- const FreqTable< 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.

- 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 89 of file support-bones.hpp.

8.41.2.2 Support() [2/3]

Constructor specifying the dimensions of the array (empty).

Definition at line 98 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 105 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 112 of file support-bones.hpp.

8.41.3 Member Function Documentation

8.41.3.1 add_counter()

Definition at line 417 of file support-meat.hpp.

8.41.3.2 add_rule() [1/2]

Definition at line 444 of file support-meat.hpp.

8.41.3.3 add_rule() [2/2]

Definition at line 454 of file support-meat.hpp.

8.41.3.4 add_rule_dyn() [1/2]

Definition at line 479 of file support-meat.hpp.

8.41.3.5 add_rule_dyn() [2/2]

Definition at line 489 of file support-meat.hpp.

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.

Definition at line 383 of file support-meat.hpp.

8.41.3.7 eval_rules_dyn()

Definition at line 514 of file support-meat.hpp.

8.41.3.8 get_counters()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >

Counters< Array_Type, Data_Counter_Type > * Support< Array_Type, Data_Counter_Type, Data_←

Rule_Type, Data_Rule_Dyn_Type >::get_counters [inline]
```

Vector of couter functions.

Definition at line 593 of file support-meat.hpp.

8.41.3.9 get_counts()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type > const std::vector< double > & Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_\leftarrow Rule_Dyn_Type >::get_counts [inline]
```

Definition at line 557 of file support-meat.hpp.

8.41.3.10 get_current_stats()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< double > * Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_\Lorentype, Dyn_Type >::get_current_stats [inline]
```

List current statistics.

Definition at line 571 of file support-meat.hpp.

8.41.3.11 get_data()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
const FreqTable< double > & Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_←
Rule_Dyn_Type >::get_data [inline]
```

Definition at line 588 of file support-meat.hpp.

8.41.3.12 get_rules()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
Rules< Array_Type, Data_Rule_Type > * Support< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_rules [inline]
```

Vector of static rules (cells to iterate).

Definition at line 598 of file support-meat.hpp.

8.41.3.13 get_rules_dyn()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >

Rules< Array_Type, Data_Rule_Dyn_Type > * Support< Array_Type, Data_Counter_Type, Data_Rule

_Type, Data_Rule_Dyn_Type >::get_rules_dyn [inline]
```

Vector of dynamic rules (to include/exclude a realizaton).

Definition at line 603 of file support-meat.hpp.

8.41.3.14 init_support()

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

8.41.3.15 print()

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

Definition at line 576 of file support-meat.hpp.

8.41.3.16 reset_array() [1/2]

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

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

8.41.3.17 reset_array() [2/2]

Definition at line 118 of file support-meat.hpp.

8.41.3.18 set_counters()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >

void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::set_

counters (

Counters< Array_Type, Data_Counter_Type > * counters_) [inline]
```

Definition at line 427 of file support-meat.hpp.

8.41.3.19 set_rules()

Definition at line 464 of file support-meat.hpp.

8.41.3.20 set_rules_dyn()

Definition at line 499 of file support-meat.hpp.

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 \( \to \) _Type >::change_stats
```

Definition at line 82 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 80 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 \cdot _n_locked
```

Definition at line 81 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 78 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 79 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 77 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 71 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 72 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 73 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 83 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 84 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 70 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 74 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 70 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 85 of file support-bones.hpp.

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

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

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.

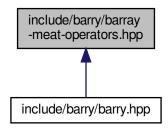
9.2 include/barry/barray-iterator.hpp File Reference

Classes

class ConstBArrayRowIter< Cell_Type, Data_Type >

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
- 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

```
#define 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

```
#define 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

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 ( ) [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.

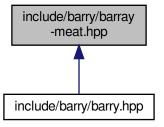
9.3.3.2 this

```
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 ROW(a) this->el_ij[a]
- #define COL(a) this->el_ji[a]

9.4.1 Macro Definition Documentation

9.4.1.1 COL

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

9.4.1.2 ROW

Definition at line 13 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:

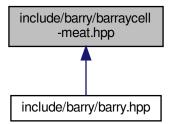


Classes

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

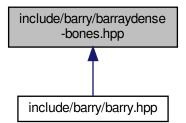
9.6 include/barry/barraycell-meat.hpp File Reference

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

```
#define 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

```
#define 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 &
```

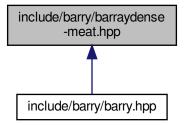
9.8.2.6 BDENSE_TYPE()

```
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 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)

9.9.1 Macro Definition Documentation

9.9.1.1 COL

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

9.9.1.2 POS

```
#define POS(  a, \\ b ) \ (b)*N + (a)
```

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

9.9.1.3 POS_N

Definition at line 26 of file barraydense-meat.hpp.

9.9.1.4 ROW

Definition at line 23 of file barraydense-meat.hpp.

9.9.1.5 ZERO_CELL

```
#define ZERO_CELL static_cast<Cell_Type>(0.0)
```

Definition at line 31 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(  a, \\ b \ ) \ (a) \ + \ (b) \ * \ \mbox{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

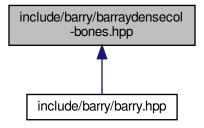
9.11.1.1 POS

```
#define POS(  a, \\ b ) (a) + (b) * dat->N
```

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.

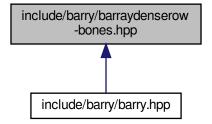
9.12.1.3 ZERO_CELL

```
#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, 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(  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]

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

```
#define BARRY_CHECK_SUPPORT(
          x,
          maxs )
```

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

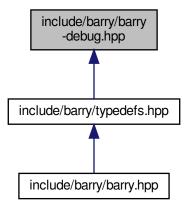
9.18.2.1 Map

```
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

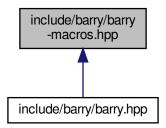
9.19.1.1 BARRY_DEBUG_LEVEL

#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);
- #define BARRY_NCORES_ARG(default) size_t ncores default

9.20.1 Macro Definition Documentation

9.20.1.1 BARRY_NCORES_ARG

Definition at line 15 of file barry-macros.hpp.

9.20.1.2 BARRY_ONE

```
#define BARRY_ONE CellCell_Type>(1.0)
```

Definition at line 7 of file barry-macros.hpp.

9.20.1.3 BARRY_ONE_DENSE

```
#define BARRY_ONE_DENSE static_cast<Cell_Type>(1.0)
```

Definition at line 8 of file barry-macros.hpp.

9.20.1.4 BARRY_UNUSED

Definition at line 10 of file barry-macros.hpp.

9.20.1.5 BARRY_ZERO

```
#define BARRY_ZERO Cell<Cell_Type>(0.0)
```

Definition at line 4 of file barry-macros.hpp.

9.20.1.6 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 29 of file barry.hpp.

9.21.1.2 BARRY_VERSION

```
#define BARRY_VERSION BARRY_VERSION_MAYOR ## . ## BARRY_VERSION_MINOR
```

Definition at line 33 of file barry.hpp.

9.21.1.3 BARRY_VERSION_MAYOR

```
#define BARRY_VERSION_MAYOR 0
```

Definition at line 31 of file barry.hpp.

9.21.1.4 BARRY_VERSION_MINOR

```
#define BARRY_VERSION_MINOR 1
```

Definition at line 32 of file barry.hpp.

9.21.1.5 COUNTER_FUNCTION

```
\begin{array}{c} \texttt{\#define COUNTER\_FUNCTION}\,(\\ & a \end{array})
```

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 92 of file barry.hpp.

9.21.1.6 COUNTER_LAMBDA

Definition at line 95 of file barry.hpp.

9.21.1.7 RULE_FUNCTION

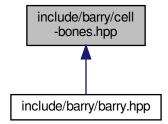
Definition at line 99 of file barry.hpp.

9.21.1.8 RULE LAMBDA

Definition at line 102 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:

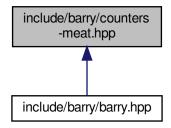


Classes

- 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>
- #define COUNTERS_TEMPLATE(a, b) template COUNTERS_TEMPLATE_ARGS() inline a COUNTERS_TYPE()

 ::b

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

- 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
- COUNTERS_TEMPLATE (COUNTERS_TYPE() &, operator=)(Counters< Array_Type
- COUNTERS TEMPLATE (void, add counter)(Counter< Array Type
- 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

Variables

- Data_Type & counter_
- Data Type &&counter noexcept
- size t i
- 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]

< 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]

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 213 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 233 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 248 of file counters-meat.hpp.

9.26.2.28 if() [2/3]

```
if ( hasher )
```

Definition at line 255 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 227 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_Type Counter_fun_type<Array_Type,Data_Type> Hasher_fun_type<Array_Type,Data_Type> Data←
Type 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.emplace_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 270 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

```
size_t i
```

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 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 265 of file counters-meat.hpp.

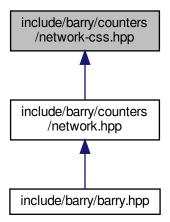
9.26.3.16 return

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_MATCH_TYPE()
- #define CSS_CASE_TRUTH() if ((i < n) && (j < n))
- #define CSS_TRUE_CELLS()
- #define CSS_CASE_PERCEIVED() else if (((i >= s) && (i < e)) & ((j >= 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_, size_t counter_type=0u)
     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_, size_t counter_type=0u)
     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_, size_t counter_type=0u)
     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_, size_t counter_type=0u)
     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 , size t counter type=0u)
     Counts mixed reciprocity errors.
• template<typename Tnet = Network>
  void counter_css_census01 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end_, size_t counter_type=0u)

    template<tvpename Tnet = Network>

  void counter_css_census02 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end , size t counter type=0u)

    template<typename Tnet = Network>

  void counter_css_census03 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end_, size_t counter_type=0u)
template<typename Tnet = Network>
  void counter_css_census04 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end_, size_t counter_type=0u)
template<typename Tnet = Network>
  void counter_css_census05 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end_, size_t counter_type=0u)
template<typename Tnet = Network>
  void counter css census06 (NetCounters< Tnet > *counters, size t netsize, const std::vector< size t >
  &end_, size_t counter_type=0u)
template<typename Tnet = Network>
  void counter_css_census07 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end_, size_t counter_type=0u)
template<typename Tnet = Network>
  void counter_css_census08 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end , size t counter type=0u)
template<typename Tnet = Network>
  void counter css census09 (NetCounters< Tnet > *counters, size t netsize, const std::vector< size t >
  &end_, size_t counter_type=0u)
template<typename Tnet = Network>
  void counter css census10 (NetCounters< Tnet > *counters, size t netsize, const std::vector< size t >
  &end , size t counter type=0u)
```

9.27.1 Macro Definition Documentation

9.27.1.1 CSS_APPEND

Definition at line 81 of file network-css.hpp.

9.27.1.2 CSS_CASE_ELSE

```
#define CSS_CASE_ELSE( )
```

Definition at line 66 of file network-css.hpp.

9.27.1.3 CSS CASE PERCEIVED

```
      \# define \ CSS\_CASE\_PERCEIVED( ) \ else \ if \ (((i >= s) \ \&\& \ (i < e)) \ \& \ ((j >= s) \ \&\& \ (j < e)))
```

Definition at line 48 of file network-css.hpp.

9.27.1.4 CSS_CASE_TRUTH

```
#define CSS_CASE_TRUTH( ) if ((i < n) && (j < n))
```

Definition at line 32 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]) \</pre>
```

throw std::logic_error("Endpoints should be specified in order.");}

Definition at line 76 of file network-css.hpp.

9.27.1.6 CSS_CHECK_SIZE_INIT

Definition at line 70 of file network-css.hpp.

9.27.1.7 CSS_MATCH_TYPE

Definition at line 22 of file network-css.hpp.

9.27.1.8 CSS_NET_COUNTER_LAMBDA_INIT

Definition at line 89 of file network-css.hpp.

9.27.1.9 CSS PERCEIVED CELLS

```
#define CSS_PERCEIVED_CELLS()

Value:
    size_t i_ = i - s; \
    size_t j_ = j - s; \
    CSS_MATCH_TYPE() \
    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 55 of file network-css.hpp.

9.27.1.10 CSS_SIZE

```
#define CSS_SIZE()

Value:
    size_t n = data.indices[0u]; \
    size_t s = data.indices[1u]; \
    size_t e = data.indices[2u]; \
    size_t ctype = data.indices[3u]; \
    size_t ego_id = data.indices[4u]; \
    if (ctype > 2) \
        throw std::range_error("Counter type should be 0, 1, or 2.");
```

Definition at line 8 of file network-css.hpp.

9.27.1.11 CSS_TRUE_CELLS

```
#define CSS_TRUE_CELLS()

Value:
    size_t i_ = i; \
    size_t j_ = j; \
    CSS_MATCH_TYPE() \
    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 39 of file network-css.hpp.

9.27.2 Function Documentation

9.27.2.1 counter_css_census01()

Definition at line 324 of file network-css.hpp.

9.27.2.2 counter_css_census02()

Definition at line 389 of file network-css.hpp.

9.27.2.3 counter_css_census03()

Definition at line 429 of file network-css.hpp.

9.27.2.4 counter css census04()

Definition at line 469 of file network-css.hpp.

9.27.2.5 counter_css_census05()

Definition at line 509 of file network-css.hpp.

9.27.2.6 counter_css_census06()

Definition at line 549 of file network-css.hpp.

9.27.2.7 counter_css_census07()

Definition at line 589 of file network-css.hpp.

9.27.2.8 counter css census08()

Definition at line 629 of file network-css.hpp.

9.27.2.9 counter_css_census09()

Definition at line 669 of file network-css.hpp.

9.27.2.10 counter_css_census10()

Definition at line 709 of file network-css.hpp.

9.27.2.11 counter_css_completely_false_recip_comiss()

Counts completely false reciprocity (comission)

Definition at line 200 of file network-css.hpp.

9.27.2.12 counter_css_completely_false_recip_omiss()

Counts completely false reciprocity (omission)

Definition at line 241 of file network-css.hpp.

9.27.2.13 counter_css_mixed_recip()

Counts mixed reciprocity errors.

Definition at line 282 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)
counter_type	Size_t indicating the type of counter to use. Possible values are: 0: Count all, 1: Only count if perceiver is involved, and 2: Only count if perceiver is not involved.

The end_ parameter should be of length ${\tt N}$ of networks - 1. It is assumed that the first network ends at netsize.

Definition at line 107 of file network-css.hpp.

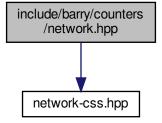
9.27.2.15 counter_css_partially_false_recip_omiss()

Counts errors of omission.

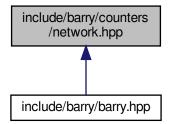
Definition at line 155 of file network-css.hpp.

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<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.
```

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

Value:

```
template<typename Tnet = Network>\
inline double (a) (const Tnet & Array, size_t i, size_t j, NetCounterData & data)
```

Function for definition of a network counter function

Definition at line 114 of file network.hpp.

9.28.1.6 NETWORK_COUNTER_LAMBDA

Value:

```
Counter_fun_type<Tnet, NetCounterData> a = \
   [](const Tnet & Array, size_t i, size_t j, NetCounterData & data)
```

Lambda function for definition of a network counter function

Definition at line 119 of file network.hpp.

9.28.1.7 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

Counter_fun_type<NetworkDense, NetCounterData> a = \
[](const NetworkDense & Array, size_t i, size_t j, NetCounterData & data)

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

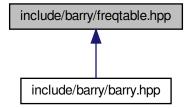
9.28.3.1 rules_zerodiag()

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:



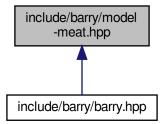
Classes

- 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:



Functions

- double update_normalizing_constant (const std::vector< double > ¶ms, const double *support, size_t k, size t n)
- double likelihood_ (const double *stats_target, const std::vector< double > ¶ms, const double normalizing_constant, size_t n_params, bool log_=false)

9.31.1 Function Documentation

9.31.1.1 likelihood_()

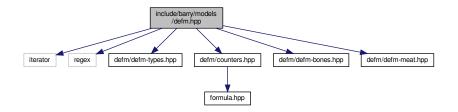
Definition at line 71 of file model-meat.hpp.

9.31.1.2 update_normalizing_constant()

Definition at line 9 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:
```

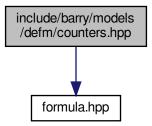


Namespaces

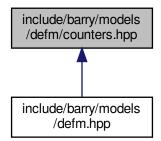
defm

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)

 Data for the counters.
- #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 >).

- $\bullet \ \ void \ rules_markov_fixed \ (DEFMRules *rules, size_t \ markov_order)\\$
- Number of edges.
 void rules_dont_become_zero (DEFMSupport *support, std::vector < size_t > ids)
 Blocks switching a one to zero.
- void rule_constrain_support (DEFMSupport *support, size_t pos, double lb, double ub)
 Overall functional gains.

9.33.1 Macro Definition Documentation

9.33.1.1 DEFM COUNTER

Function for definition of a network counter function

Definition at line 47 of file counters.hpp.

9.33.1.2 DEFM_COUNTER_LAMBDA

[](const DEFMArray & Array, size_t i, size_t j, DEFMCounterData & data) -> double

Lambda function for definition of a network counter function

barry::Counter_fun_type<DEFMArray, DEFMCounterData> a = \

Definition at line 51 of file counters.hpp.

9.33.1.3 DEFM_RULE

Function for definition of a network counter function

Definition at line 61 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 65 of file counters.hpp.

9.33.1.5 DEFM_RULEDYN_LAMBDA

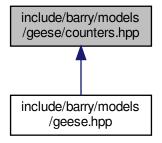
Lambda function for definition of a network counter function

Definition at line 71 of file counters.hpp.

9.33.1.6 UNI_SUB

9.34 include/barry/models/geese/counters.hpp File Reference

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



Functions

- void rule_leafs (PhyloSupport *support)
- 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_RULE_LAMBDA(a)

Extension of a simple counter.

- #define PHYLO COUNTER LAMBDA(a)
- #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)

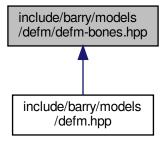
Used when all the functions are in 0 (like the root node prob.)

• 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.)

9.35 include/barry/models/defm/defm-bones.hpp File Reference

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

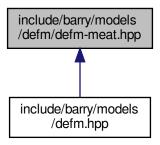


Classes

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 36 of file defm-meat.hpp.

9.36.1.2 DEFM_RANGES

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

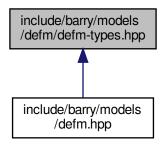
9.36.2 Function Documentation

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
- $\bullet \ \ type def \ barry:: Rule < DEFMArray, \ DEFMRule Data > DEFMRule \\$
- typedef barry::Rules < DEFMArray, DEFMRuleData > DEFMRules
- typedef barry::Rule < DEFMArray, DEFMRuleDynData > DEFMRuleDyn
- typedef barry::Rules < DEFMArray, DEFMRuleDynData > DEFMRulesDyn

9.37.1 Typedef Documentation

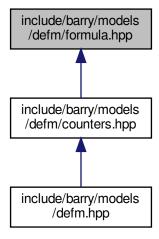
9.37.1.1 **DEFMArray**

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, std::string &covar_name, std::string &vname)

Parses a motif formula.

9.38.1 Function Documentation

9.38.1.1 defm_motif_parser()

```
void defm_motif_parser (
    std::string formula,
    std::vector< size_t > & locations,
    std::vector< bool > & signs,
    size_t m_order,
    size_t y_ncol,
    std::string & covar_name,
    std::string & vname ) [inline]
```

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 preceeding 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.

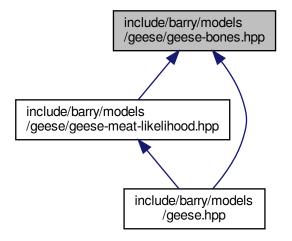
9.41 include/barry/models/geese/flock-meat.hpp File Reference

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 71 of file geese-bones.hpp.

9.42.2.4 vector_caster()

```
template<typename Ta , typename Tb >  $ $ std::vector< Ta > vector\_caster ( \\ const std::vector< Tb > & x ) [inline]
```

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

9.43 include/barry/models/geese/geese-meat-constructors.hpp File Reference

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

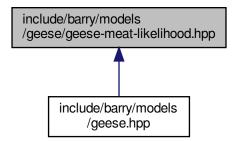


9.44 include/barry/models/geese/geese-meat-likelihood.hpp File Reference

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



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



Functions

void pset_loop (size_t n, size_t s, size_t nfunctions, const size_t node_id, const size_t array_id, std::vector< double > &totprob_n, const std::vector< double > &par0, const std::vector< std::vector< bool >> &states, const std::vector< PhyloArray > &psets, const std::vector< std::vector< size_t >> &locations, const std::vector< geese::Node *> &node_offspring, const double *psetprobs)

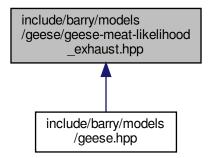
9.44.1 Function Documentation

9.44.1.1 pset_loop()

```
void pset_loop (
    size_t n,
    size_t s,
    size_t nfunctions,
    const size_t node_id,
    const size_t array_id,
    std::vector< double > & totprob_n,
    const std::vector< double > & par0,
    const std::vector< std::vector< bool >> & states,
    const std::vector< PhyloArray > & psets,
    const std::vector< std::vector< size_t > > & locations,
    const std::vector< geese::Node * > & node_offspring,
    const double * psetprobs ) [inline]
```

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

9.45 include/barry/models/geese/geese-meat-likelihood_exhaust.hpp File Reference



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

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



9.47 include/barry/models/geese/geese-meat-predict_exhaust.hpp File Reference

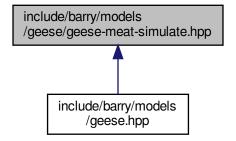


9.48 include/barry/models/geese/geese-meat-predict_sim.hpp File Reference

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



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



9.50 include/barry/models/geese/geese-meat.hpp File Reference

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



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

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



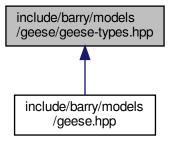
Classes

• 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

Macros

• #define POS(a, b) (b)*N + (a)

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 Macro Definition Documentation

9.52.1.1 POS

```
#define POS(  a, \\ b ) (b)*N + (a)
```

Definition at line 4 of file geese-types.hpp.

9.52.2 Typedef Documentation

9.52.2.1 PhyloArray

```
typedef barry::BArrayDense<size_t, NodeData> PhyloArray
```

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

9.52.2.2 PhyloCounter

```
typedef barry::Counter<PhyloArray, PhyloCounterData > PhyloCounter
```

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

9.52.2.3 PhyloCounters

```
{\tt typedef\ barry::Counters} < {\tt PhyloArray,\ PhyloCounterData} > {\tt PhyloCounters}
```

Definition at line 105 of file geese-types.hpp.

9.52.2.4 PhyloModel

```
{\tt typedef\ barry::Model < PhyloArray,\ PhyloCounterData,\ PhyloRuleData,\ PhyloRuleDynData > PhyloModel}
```

Definition at line 115 of file geese-types.hpp.

9.52.2.5 PhyloPowerSet

typedef barry::PowerSet<PhyloArray, PhyloRuleData> PhyloPowerSet

Definition at line 116 of file geese-types.hpp.

9.52.2.6 PhyloRule

typedef barry::Rule<PhyloArray,PhyloRuleData> PhyloRule

Definition at line 107 of file geese-types.hpp.

9.52.2.7 PhyloRuleData

typedef std::vector< std::pair< size_t, size_t > > PhyloRuleData

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

9.52.2.8 PhyloRuleDyn

typedef barry::Rule<PhyloArray,PhyloRuleDynData> PhyloRuleDyn

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

9.52.2.9 PhyloRules

typedef barry::Rules<PhyloArray,PhyloRuleData> PhyloRules

Definition at line 108 of file geese-types.hpp.

9.52.2.10 PhyloRulesDyn

typedef barry::Rules<PhyloArray,PhyloRuleDynData> PhyloRulesDyn

Definition at line 111 of file geese-types.hpp.

9.52.2.11 PhyloStatsCounter

typedef barry::StatsCounter<PhyloArray, PhyloCounterData> PhyloStatsCounter

Definition at line 114 of file geese-types.hpp.

9.52.2.12 PhyloSupport

typedef barry::Support<PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloSupport

Definition at line 113 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.
- $\bullet \ \ {\it class Rules} {< Array_Type, Data_Type} >$

Vector of objects of class Rule.

Functions

template<typename Array_Type , typename Data_Type >
 bool rule_fun_default (const Array_Type *array, size_t i, size_t j, Data_Type *dat)

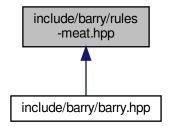
9.56.1 Function Documentation

9.56.1.1 rule_fun_default()

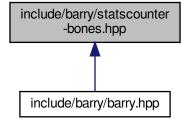
Definition at line 5 of file rules-bones.hpp.

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
- 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_t i
- STATSCOUNTER_TEMPLATE (std::vector< std::string >, get_names)() const
- STATSCOUNTER_TEMPLATE (std::vector< std::string >, get_descriptions)() const

Variables

```
Data_Type & counterEmptyArray = *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_
- size_t j

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_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!")
```

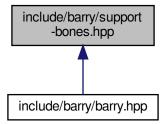
Definition at line 66 of file statscounter-meat.hpp.

9.59.3.9 return

return

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

9.60 include/barry/support-bones.hpp File Reference

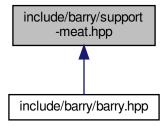


Classes

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

9.61.1 Macro Definition Documentation

9.61.1.1 BARRY_SUPPORT_MEAT_HPP

#define BARRY_SUPPORT_MEAT_HPP 1

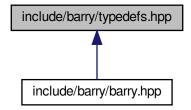
Definition at line 2 of file support-meat.hpp.

9.62 include/barry/typedefs.hpp File Reference

```
#include "barry-configuration.hpp"
#include "barry-debug.hpp"
#include "progress.hpp"
Include dependency graph for typedefs.hpp:
```



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



Classes

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

Namespaces

- CHECK
 - Integer constants used to specify which cell should be check.
- EXISTS

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

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, bata_Type &)>
          Counter and rule functions.
    • template<typename Array_Type , typename Data_Type >
      using Rule_fun_type = std::function< bool(const Array_Type &, size_t, 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
```

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.

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.

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