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. The goal of the library 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.

This 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."

Examples

Counting statistics in a graph

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

```
#include <iostream>
#include <ostream>
#include "../include/barry.hpp"
typedef std::vector< unsigned int > vuint;
int main() {
```

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```
// Creating network of size six with five ties
  netcounters::Network net(
       6, 6,
      {0, 0, 4, 4, 2, 0, 1},
      {1, 2, 0, 2, 4, 0, 1}
  // How does this looks like?
  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();
  std::cout «
                         : " « counts[0] « std::endl «
    "Transitive triads : " « counts[U] « std::endl «
"Transitive triads : " « counts[1] « std::endl «
"Isolates : " « counts[2] « std::endl «
                         : " « counts[3] « std::endl « : " « counts[4] « std::endl;
    "C triads
    "Mutuals
  return 0;
Compiling this program using g++
g++ -std=c++11 -Wall -pedantic 08-counts.cpp -o counts && ./counts
Yields the following output:
Current view
        1
               1
   1,]
            1
   2,] .
                   . 1
   3,] . . . . . . 4,] 1 . 1 .
   5,]
  0,] .
1,] 1
           1
               1
  2,] 1 .
                      1
   3,] . . .
4,] 1 . 1
   5,] . . . .
Edges
Transitive triads : 3
Isolates
C triads
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:

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EFMArray counters	12
nylo counters	2(
nylo rules	2

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Chapter 3

Class Index

3.1 Class List

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

BArray< Cell_Type, Data_Type >
Baseline class for binary arrays
BArrayCell < Cell_Type, Data_Type >
BArrayCell_const< Cell_Type, Data_Type >
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 86
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
DEFMCounterData
Data class used to store arbitrary uint or double vectors
DEFMData
Data class for DEFM arrays
Entries < Cell_Type >
A wrapper class to store source, target, val from a BArray object 118
Flock
A Flock is a group of Geese
FreqTable < T >
Frequency table of vectors

8 Class Index

Geese	
Annotated Phylo Model	128
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:	147
NetCounterData	
Data class used to store arbitrary uint or double vectors	162
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A single node for the model	165
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Data definition for the PhyloArray class	171
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PowerSet < Array_Type, Data_Rule_Type >	
Powerset of a binary array	177
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Rule < Array_Type, Data_Type >	
Rule for determining if a cell should be included in a sequence	185
Rules < Array_Type, Data_Type >	
Vector of objects of class Rule	187
StatsCounter< Array_Type, Data_Type >	
Count stats for a single Array	191
Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >	
Compute the support of sufficient statistics	194
vecHasher< T >	

Chapter 4

File Index

4.1 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/defm.hpp
include/barry/counters/network-css.hpp
include/barry/counters/network.hpp
include/barry/counters/phylo.hpp
include/barry/models/defm.hpp
include/barry/models/geese.hpp
include/barry/models/defm/defm-bones.hpp
include/barry/models/defm/defm-meat.hpp
include/barry/models/geese/flock-bones.hpp
include/barry/models/geese/flock-meat.hpp
include/barry/models/geese/geese-bones.hpp
include/barry/models/geese/geese-meat-constructors.hpp
include/barry/models/geese/geese-meat-likelihood.hpp
include/barry/models/geese/geese-meat-likelihood_exhaust.hpp
include/barry/models/geese/geese-meat-predict.hpp
include/barry/models/geese/geese-meat-predict_exhaust.hpp
include/barry/models/geese/geese-meat-predict_sim.hpp
include/barry/models/geese/geese-meat-simulate.hpp
include/barry/models/geese/geese-meat.hpp
include/barry/models/geese/geese-node-bones.hpp

Chapter 5

Module Documentation

5.1 Counting

Classes

class DEFMData

Data class for DEFM arrays.

class NetworkData

Data class for Networks.

class NodeData

Data definition for the PhyloArray class.

class Counter< Array_Type, Data_Type >

A counter function based on change statistics.

5.1.1 Detailed Description

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

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

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

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

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

5.2 Statistical Models

Statistical models available in barry.

Classes

class Model < Array_Type, Data_Counter_Type, Data_Rule_Type, 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 Geese

Annotated Phylo Model.

class Flock

A Flock is a group of Geese.

5.2.1 Detailed Description

Statistical models available in barry.

5.3 **DEFMArray** counters

Counters for network models.

Functions

```
    template < typename Tarray = DEFMArray > void counter_ones (DEFMCounters < Tarray > *counters, int covar_index=-1)
```

Prevalence of ones.

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, uint 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, uint attr_id, double alpha=1.0, double tail_head=true)
     Sum of attribute difference between ego and alter to pow(alpha)

    NETWORK COUNTER (init single attr)

• template<typename Tnet = Network>
  void counter_nodeicov (NetCounters < Tnet > *counters, uint attr_id)
• template<typename Tnet = Network>
  void counter_nodeocov (NetCounters< Tnet > *counters, uint attr_id)
• template<typename Tnet = Network>
  void counter_nodecov (NetCounters< Tnet > *counters, uint attr_id)
• template<typename Tnet = Network>
  void counter nodematch (NetCounters < Tnet > *counters, uint attr id)
• template<typename Tnet = Network>
  void counter_idegree (NetCounters< Tnet > *counters, std::vector< uint > d)
     Counts number of vertices with a given in-degree.

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

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

    template<> void counter odegree (NetCounters< NetworkDense > *counters, std::vector< uint > d)

• template<typename Tnet = Network>
  void\ counter\_degree\ (NetCounters < Tnet > *counters,\ std::vector < uint > d)
     Counts number of vertices with a given out-degree.
```

5.3.1 Detailed Description

Counters for network models.

Parameters

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

5.3.2 Function Documentation

5.3.2.1 counter_absdiff()

```
uint attr_id,
double alpha = 1.0 ) [inline]
```

Sum of absolute attribute difference between ego and alter.

Definition at line 910 of file network.hpp.

5.3.2.2 counter_ctriads() [1/2]

Definition at line 665 of file network.hpp.

5.3.2.3 counter_ctriads() [2/2]

Definition at line 610 of file network.hpp.

5.3.2.4 counter_degree()

Counts number of vertices with a given out-degree.

Definition at line 1328 of file network.hpp.

5.3.2.5 counter_density()

Definition at line 731 of file network.hpp.

5.3.2.6 counter_diff()

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

Definition at line 955 of file network.hpp.

5.3.2.7 counter_edges()

Number of edges.

Definition at line 152 of file network.hpp.

5.3.2.8 counter_idegree() [1/2]

Definition at line 1172 of file network.hpp.

5.3.2.9 counter_idegree() [2/2]

Counts number of vertices with a given in-degree.

Definition at line 1125 of file network.hpp.

5.3.2.10 counter_idegree15() [1/2]

Definition at line 787 of file network.hpp.

5.3.2.11 counter_idegree15() [2/2]

Definition at line 759 of file network.hpp.

5.3.2.12 counter_isolates() [1/2]

Definition at line 215 of file network.hpp.

5.3.2.13 counter_isolates() [2/2]

Number of isolated vertices.

Definition at line 175 of file network.hpp.

5.3.2.14 counter_istar2() [1/2]

Definition at line 338 of file network.hpp.

5.3.2.15 counter_istar2() [2/2]

Definition at line 312 of file network.hpp.

5.3.2.16 counter_mutual()

Number of mutual ties.

Definition at line 256 of file network.hpp.

5.3.2.17 counter_nodecov()

Definition at line 1068 of file network.hpp.

5.3.2.18 counter_nodeicov()

Definition at line 1018 of file network.hpp.

5.3.2.19 counter_nodematch()

Definition at line 1093 of file network.hpp.

5.3.2.20 counter_nodeocov()

Definition at line 1043 of file network.hpp.

5.3.2.21 counter_odegree() [1/2]

Definition at line 1273 of file network.hpp.

5.3.2.22 counter_odegree() [2/2]

Counts number of vertices with a given out-degree.

Definition at line 1225 of file network.hpp.

5.3.2.23 counter_odegree15() [1/2]

Definition at line 864 of file network.hpp.

5.3.2.24 counter_odegree15() [2/2]

Definition at line 836 of file network.hpp.

5.3.2.25 counter_ones()

Prevalence of ones.

Canonical term.

Template Parameters

Tarray	

Parameters

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

Definition at line 167 of file defm.hpp.

5.3.2.26 counter_ostar2() [1/2]

Definition at line 404 of file network.hpp.

5.3.2.27 counter_ostar2() [2/2]

Definition at line 376 of file network.hpp.

5.3.2.28 counter_ttriads() [1/2]

Definition at line 531 of file network.hpp.

5.3.2.29 counter_ttriads() [2/2]

Definition at line 441 of file network.hpp.

5.3.2.30 NETWORK_COUNTER()

Definition at line 999 of file network.hpp.

5.4 Phylo counters

Counters for phylogenetic modeling.

Functions

- void counter_overall_gains (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)
 Overall functional gains.
- void counter_gains (PhyloCounters *counters, std::vector< uint > nfun, unsigned int duplication=DEFAULT_DUPLICATION)

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

k genes gain function nfun

- void counter_genes_changing (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)
- void counter_preserve_pseudogene (PhyloCounters *counters, unsigned int nfunA, unsigned int nfunB, unsigned int duplication=DEFAULT DUPLICATION)

Keeps track of how many pairs of genes preserve pseudostate.

- void counter_prop_genes_changing (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)
- void counter_overall_loss (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)
 Overall functional loss.
- void counter_maxfuns (PhyloCounters *counters, uint lb, uint ub, unsigned int duplication=DEFAULT_DUPLICATION)

 Cap the number of functions per gene.
- void counter_loss (PhyloCounters *counters, std::vector< uint > nfun, unsigned int duplication=DEFAULT_DUPLICATION)

 Total count of losses for an specific function.
- void counter_overall_changes (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Total number of changes. Use this statistic to account for "preservation".
- void counter_subfun (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

 Total count of Sub-functionalization events.
- void counter_cogain (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

5.4 Phylo counters 21

Co-evolution (joint gain or loss)

void counter_longest (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

Longest branch mutates (either by gain or by loss)

• void counter_neofun (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

Total number of neofunctionalization events.

void counter_pairwise_neofun_singlefun (PhyloCounters *counters, uint nfunA, unsigned int duplication=DEFAULT_DUPLICATI
 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, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION Total number of neofunctionalization events.
- void counter_co_opt (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

 Function co-opting.
- void counter_k_genes_changing (PhyloCounters *counters, unsigned int k, unsigned int duplication=DEFAULT_DUPLICATION)
 Indicator function. Equals to one if k genes changed and zero otherwise.
- void counter_less_than_p_prop_genes_changing (PhyloCounters *counters, double p, unsigned int duplication=DEFAULT DUPLICATION)

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

- void counter_gains_from_0 (PhyloCounters *counters, std::vector< uint > nfun, unsigned int duplication=DEFAULT_DUPLICAT

 Used when all the functions are in 0 (like the root node prob.)
- void counter_overall_gains_from_0 (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Used when all the functions are in 0 (like the root node prob.)
- void counter_pairwise_overall_change (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Used when all the functions are in 0 (like the root node prob.)
- void counter_pairwise_preserving (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT DUPLICATION)

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

 void counter_pairwise_first_gain (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

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

5.4.1 Detailed Description

Counters for phylogenetic modeling.

Parameters

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

5.4.2 Function Documentation

5.4.2.1 counter_co_opt()

```
uint nfunB,
unsigned int duplication = DEFAULT_DUPLICATION ) [inline]
```

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:

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

This algorithm implements the change statistic.

Definition at line 1393 of file phylo.hpp.

5.4.2.2 counter_cogain()

Co-evolution (joint gain or loss)

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

Definition at line 888 of file phylo.hpp.

5.4.2.3 counter_gains()

Functional gains for a specific function (nfun).

Definition at line 193 of file phylo.hpp.

5.4.2.4 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 1727 of file phylo.hpp.

5.4 Phylo counters 23

5.4.2.5 counter_gains_k_offspring()

k genes gain function nfun

Definition at line 253 of file phylo.hpp.

5.4.2.6 counter_genes_changing()

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

Definition at line 325 of file phylo.hpp.

5.4.2.7 counter_k_genes_changing()

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

Definition at line 1491 of file phylo.hpp.

5.4.2.8 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 1611 of file phylo.hpp.

5.4.2.9 counter_longest()

Longest branch mutates (either by gain or by loss)

Definition at line 945 of file phylo.hpp.

5.4.2.10 counter_loss()

Total count of losses for an specific function.

Definition at line 688 of file phylo.hpp.

5.4.2.11 counter_maxfuns()

Cap the number of functions per gene.

Definition at line 626 of file phylo.hpp.

5.4.2.12 counter_neofun()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1115 of file phylo.hpp.

5.4 Phylo counters 25

5.4.2.13 counter_neofun_a2b()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1260 of file phylo.hpp.

5.4.2.14 counter_overall_changes()

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

Definition at line 740 of file phylo.hpp.

5.4.2.15 counter_overall_gains()

Overall functional gains.

Total number of gains (irrespective of the function).

Definition at line 155 of file phylo.hpp.

5.4.2.16 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 1793 of file phylo.hpp.

5.4.2.17 counter_overall_loss()

Overall functional loss.

Definition at line 578 of file phylo.hpp.

5.4.2.18 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 2045 of file phylo.hpp.

5.4.2.19 counter_pairwise_neofun_singlefun()

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

Definition at line 1196 of file phylo.hpp.

5.4.2.20 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 1841 of file phylo.hpp.

5.4 Phylo counters 27

5.4.2.21 counter_pairwise_preserving()

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

Needs to specify function a. sum $x(a)^3(1-x(b))^3 + x(b)^3(1-x(a))^3 + x(a)^3 + x(b)^3 + (1-x(a))^3 + (1-x(b))^3$ Definition at line 1906 of file phylo.hpp.

5.4.2.22 counter_preserve_pseudogene()

Keeps track of how many pairs of genes preserve pseudostate.

Definition at line 394 of file phylo.hpp.

5.4.2.23 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 476 of file phylo.hpp.

5.4.2.24 counter_subfun()

Total count of Sub-functionalization events.

It requires to specify data = {funA, funB}

Definition at line 799 of file phylo.hpp.

5.5 Phylo rules

Rules for phylogenetic modeling.

Classes

• class PhyloRuleDynData

Functions

void rule_dyn_limit_changes (PhyloSupport *support, uint pos, uint lb, uint ub, unsigned int duplication=DEFAULT_DUPLICATIC
 Overall functional gains.

5.5.1 Detailed Description

Rules for phylogenetic modeling.

Parameters

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

5.5.2 Function Documentation

5.5.2.1 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

5.5 Phylo rules 29

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

Definition at line 2177 of file phylo.hpp.

Chapter 6

Namespace Documentation

6.1 barry Namespace Reference

barry: Your go-to motif accountant

Namespaces

counters

Tree class and Treelterator class.

6.1.1 Detailed Description

barry: Your go-to motif accountant

6.2 barry::counters Namespace Reference

Tree class and Treelterator class.

Namespaces

- network
- phylo

6.2.1 Detailed Description

Tree class and Treelterator class.

6.3 barry::counters::network Namespace Reference

6.4 barry::counters::phylo Namespace Reference

6.5 CHECK Namespace Reference

Integer constants used to specify which cell should be check.

Variables

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

6.5.1 Detailed Description

Integer constants used to specify which cell should be check.

6.5.2 Variable Documentation

6.5.2.1 BOTH

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

Definition at line 28 of file typedefs.hpp.

6.5.2.2 NONE

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

Definition at line 29 of file typedefs.hpp.

6.5.2.3 ONE

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

Definition at line 30 of file typedefs.hpp.

6.5.2.4 TWO

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

Definition at line 31 of file typedefs.hpp.

6.6 EXISTS Namespace Reference

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

Variables

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

6.6.1 Detailed Description

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

6.6.2 Variable Documentation

6.6.2.1 AS_ONE

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

Definition at line 46 of file typedefs.hpp.

6.6.2.2 AS_ZERO

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

Definition at line 45 of file typedefs.hpp.

6.6.2.3 BOTH

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

Definition at line 39 of file typedefs.hpp.

6.6.2.4 NONE

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

Definition at line 40 of file typedefs.hpp.

6.6.2.5 ONE

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

Definition at line 41 of file typedefs.hpp.

6.6.2.6 TWO

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

Definition at line 42 of file typedefs.hpp.

6.6.2.7 UKNOWN

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

Definition at line 44 of file typedefs.hpp.

Chapter 7

Class Documentation

7.1 BArray < Cell_Type, Data_Type > Class Template Reference

Baseline class for binary arrays.

#include <barray-bones.hpp>

Public Member Functions

- bool operator== (const BArray< Cell_Type, Data_Type > &Array_)
- ∼BArray ()
- void out_of_range (uint i, uint j) const
- Cell_Type get_cell (uint i, uint j, bool check_bounds=true) const
- std::vector< Cell_Type > get_col_vec (uint i, bool check_bounds=true) const
- std::vector< Cell Type > get row vec (uint i, bool check bounds=true) const
- void get_col_vec (std::vector< Cell_Type > *x, uint i, bool check_bounds=true) const
- void get_row_vec (std::vector< Cell_Type > *x, uint i, bool check_bounds=true) const
- const Row_type< Cell_Type > & row (uint i, bool check_bounds=true) const
- const Col_type< Cell_Type > & col (uint i, bool check_bounds=true) const
- Entries < Cell_Type > get_entries () const

Get the edgelist.

- void transpose ()
- void clear (bool hard=true)
- void resize (uint N_, uint M_)
- void reserve ()
- void print (const 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 unsigned int vector ranging from 0 to M_
target	When true tries to add repeated observations.

Generated by Doxygen

• BArray ()

Zero-size array.

BArray (uint N_, uint M_)

Empty array.

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

Edgelist with data.

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

Edgelist with no data (simpler)

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

Move operator.

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

void set_data (Data_Type *data_, bool delete_data_=false)

Set the data object.

- 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 (uint i, uint j, bool check_bounds=true) const
- uint nrow () const noexcept
- uint ncol () const noexcept
- uint nnozero () const noexcept
- Cell< Cell_Type > default_val () const

Cell-wise insertion/deletion

Parameters

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

BArray< Cell_Type, Data_Type > & operator+= (const std::pair< uint, uint > &coords)

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

Column/row wise interchange

- void swap rows (uint i0, uint i1, bool check bounds=true)
- void swap_cols (uint j0, uint j1, bool check bounds=true)
- void zero row (uint i, bool check bounds=true)
- void zero col (uint j, bool check bounds=true)

Arithmetic operators

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

Public Attributes

bool visited = false

Friends

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

7.1.1 Detailed Description

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

Baseline class for binary arrays.

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

Template Parameters

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

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

7.1.2 Constructor & Destructor Documentation

7.1.2.1 BArray() [1/6]

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

Zero-size array.

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

7.1.2.2 BArray() [2/6]

Empty array.

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

7.1.2.3 BArray() [3/6]

Edgelist with data.

7.1.2.4 BArray() [4/6]

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

Edgelist with no data (simpler)

7.1.2.5 BArray() [5/6]

Copy constructor.

7.1.2.6 BArray() [6/6]

Move operator.

7.1.2.7 \sim BArray()

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

7.1.3 Member Function Documentation

7.1.3.1 clear()

7.1.3.2 col()

7.1.3.3 D() [1/2]

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

7.1.3.4 D() [2/2]

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

7.1.3.5 default_val()

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

7.1.3.6 flush_data()

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

7.1.3.7 get_cell()

7.1.3.8 get_col_vec() [1/2]

7.1.3.9 get_col_vec() [2/2]

7.1.3.10 get entries()

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

Get the edgelist.

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

Returns

Entries<Cell_Type>

7.1.3.11 get_row_vec() [1/2]

7.1.3.12 get_row_vec() [2/2]

```
template<typename Cell_Type = bool, typename Data_Type = bool>
std::vector< Cell_Type > BArray< Cell_Type, Data_Type >::get_row_vec (
    uint i,
    bool check_bounds = true) const
```

7.1.3.13 insert_cell() [1/3]

7.1.3.14 insert_cell() [2/3]

7.1.3.15 insert_cell() [3/3]

7.1.3.16 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 232 of file barray-bones.hpp.

7.1.3.17 is_empty()

7.1.3.18 ncol()

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

7.1.3.19 nnozero()

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

7.1.3.20 nrow()

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

7.1.3.21 operator()() [1/2]

7.1.3.22 operator()() [2/2]

7.1.3.23 operator*=()

7.1.3.24 operator+=() [1/3]

7.1.3.25 operator+=() [2/3]

7.1.3.26 operator+=() [3/3]

7.1.3.27 operator-=() [1/3]

7.1.3.28 operator-=() [2/3]

7.1.3.29 operator-=() [3/3]

7.1.3.30 operator/=()

7.1.3.31 operator=() [1/2]

Move assignment.

7.1.3.32 operator=() [2/2]

Assignment constructor.

7.1.3.33 operator==()

7.1.3.34 out_of_range()

7.1.3.35 print()

7.1.3.36 reserve()

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

7.1.3.37 resize()

7.1.3.38 rm_cell()

7.1.3.39 row()

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

7.1.3.40 set_data()

Set the data object.

Parameters

data_	
delete_ <i>←</i>	
data_	

7.1.3.41 swap cells()

7.1.3.42 swap_cols()

7.1.3.43 swap_rows()

7.1.3.44 toggle_cell()

7.1.3.45 toggle_lock()

7.1.3.46 transpose()

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

7.1.3.47 zero_col()

7.1.3.48 zero_row()

7.1.4 Friends And Related Function Documentation

7.1.4.1 BArrayCell< Cell_Type, Data_Type >

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

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

7.1.4.2 BArrayCell_const < Cell_Type, Data_Type >

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

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

7.1.5 Member Data Documentation

7.1.5.1 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 48 of file barray-bones.hpp.

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

• include/barry/barray-bones.hpp

7.2 BArrayCell< Cell_Type, Data_Type > Class Template Reference

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

Public Member Functions

- BArrayCell (BArray < Cell_Type, Data_Type > *Array_, uint i_, uint j_, bool check_bounds=true)
- ∼BArrayCell ()
- void operator= (const Cell_Type &val)
- void operator+= (const Cell_Type &val)
- void operator-= (const Cell_Type &val)
- void operator*= (const Cell_Type &val)
- void operator/= (const Cell_Type &val)
- operator Cell_Type () const
- bool operator== (const Cell_Type &val) const

7.2.1 Detailed Description

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

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

7.2.2 Constructor & Destructor Documentation

7.2.2.1 BArrayCell()

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

7.2.2.2 ∼BArrayCell()

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

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

7.2.3 Member Function Documentation

7.2.3.1 operator Cell_Type()

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

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

7.2.3.2 operator*=()

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

7.2.3.3 operator+=()

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

7.2.3.4 operator-=()

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

7.2.3.5 operator/=()

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

7.2.3.6 operator=()

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

7.2.3.7 operator==()

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

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

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

7.3 BArrayCell_const< Cell_Type, Data_Type > Class Template Reference

#include <barraycell-bones.hpp>

Public Member Functions

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

7.3.1 Detailed Description

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

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

7.3.2 Constructor & Destructor Documentation

7.3.2.1 BArrayCell const()

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

7.3.2.2 ~BArrayCell_const()

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

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

7.3.3 Member Function Documentation

7.3.3.1 operator Cell_Type()

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

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

7.3.3.2 operator"!=()

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

7.3.3.3 operator<()

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

7.3.3.4 operator<=()

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

7.3.3.5 operator==()

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

7.3.3.6 operator>()

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

7.3.3.7 operator>=()

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

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

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

7.4 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 (uint i, uint j) const
- Cell_Type get_cell (uint i, uint j, bool check_bounds=true) const
- std::vector< Cell_Type > get_col_vec (uint i, bool check_bounds=true) const
- std::vector< Cell_Type > get_row_vec (uint i, bool check_bounds=true) const
- void get_col_vec (std::vector< Cell_Type > *x, uint i, bool check_bounds=true) const
- void get_row_vec (std::vector< Cell_Type > *x, uint i, bool check_bounds=true) const
- BArrayDenseRow< Cell_Type, Data_Type > & row (uint i, bool check_bounds=true)
- const BArrayDenseRow_const< Cell_Type, Data_Type > row (uint i, bool check_bounds=true) const
- BArrayDenseCol< Cell_Type, Data_Type > & col (uint j, bool check_bounds=true)
- const BArrayDenseCol_const< Cell_Type, Data_Type > col (uint j, bool check_bounds=true) const
- Entries < Cell_Type > get_entries () const

Get the edgelist.

- void transpose ()
- void clear (bool hard=true)
- void resize (uint N , uint M)
- · void reserve ()
- void print (const char *fmt=nullptr,...) const
- · bool is dense () const noexcept
- const std::vector< Cell_Type > & get_data () const
- const Cell_Type rowsum (unsigned int i) const
- const Cell_Type colsum (unsigned int i) const

Constructors

Parameters

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

• BArrayDense ()

Zero-size array.

BArrayDense (uint N_, uint M_)

Empty array.

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

Edgelist with data.

BArrayDense (uint N_, uint M_, const std::vector< uint > &source, const std::vector< uint > &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
 - Move operator.
- BArrayDense< Cell_Type, Data_Type > & operator= (BArrayDense< Cell_Type, Data_Type > &&x)
 noexcept

Move assignment.

void set_data (Data_Type *data_, bool delete_data_=false)

Set the data object.

- Data_Type * D ()
- 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 (uint i, uint j, bool check_bounds=true) const
- · uint nrow () const noexcept
- uint ncol () const noexcept
- · uint 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< uint, uint > &coords)
- BArrayDense< Cell_Type, Data_Type > & operator-= (const std::pair< uint, uint > &coords)
 BArrayDenseCell< Cell_Type, Data_Type > operator() (uint i, uint j, bool check_bounds=true)
- const Cell_Type operator() (uint i, uint j, bool check_bounds=true) const
- void rm cell (uint i, uint j, bool check bounds=true, bool check exists=true)
- void insert_cell (uint i, uint j, const Cell< Cell_Type > &v, bool check_bounds, bool check exists)
- void insert cell (uint i, uint j, Cell Type v, bool check bounds, bool check exists)
- void swap_cells (uint i0, uint j0, uint i1, uint j1, bool check_bounds=true, int check_exists=CHECK::BOTH, int *report=nullptr)
- void toggle_cell (uint i, uint j, bool check_bounds=true, int check_exists=EXISTS::UKNOWN)
- void toggle_lock (uint i, uint j, bool check_bounds=true)

Column/row wise interchange

- void swap_rows (uint i0, uint i1, bool check_bounds=true)
- void swap cols (uint j0, uint j1, bool check bounds=true)
- void zero_row (uint i, bool check_bounds=true)
- void zero col (uint j, bool check bounds=true)

Arithmetic operators

- BArrayDense< Cell_Type, Data_Type > & operator+= (const BArrayDense< Cell_Type, Data_Type >
- BArrayDense < Cell_Type, Data_Type > & operator+= (const Cell_Type &rhs)
- BArrayDense < Cell Type, Data Type > & operator = (const BArrayDense < Cell Type, Data Type >
- 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 >

7.4.1 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 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<unsigned int, Cell Type> >.

Template Parameters

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

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

7.4.2 Constructor & Destructor Documentation

7.4.2.1 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.

7.4.2.2 BArrayDense() [2/6]

Empty array.

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

7.4.2.3 BArrayDense() [3/6]

Edgelist with data.

7.4.2.4 BArrayDense() [4/6]

Edgelist with no data (simpler)

7.4.2.5 BArrayDense() [5/6]

Copy constructor.

7.4.2.6 BArrayDense() [6/6]

Move operator.

7.4.2.7 ∼BArrayDense()

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

7.4.3 Member Function Documentation

7.4.3.1 clear()

7.4.3.2 col() [1/2]

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

7.4.3.3 col() [2/2]

Definition at line 468 of file barraydense-meat.hpp.

7.4.3.4 colsum()

7.4.3.5 D() [1/2]

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

7.4.3.6 D() [2/2]

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

7.4.3.7 default_val()

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

7.4.3.8 get_cell()

7.4.3.9 get_col_vec() [1/2]

7.4.3.10 get_col_vec() [2/2]

7.4.3.11 get_data()

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

7.4.3.12 get_entries()

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

Get the edgelist.

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

Returns

Entries<Cell_Type>

7.4.3.13 get_row_vec() [1/2]

7.4.3.14 get_row_vec() [2/2]

```
template<typename Cell_Type = bool, typename Data_Type = bool>
std::vector< Cell_Type > BArrayDense< Cell_Type, Data_Type >::get_row_vec (
    uint i,
    bool check_bounds = true ) const
```

7.4.3.15 insert_cell() [1/2]

7.4.3.16 insert cell() [2/2]

7.4.3.17 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 254 of file barraydense-bones.hpp.

7.4.3.18 is_empty()

7.4.3.19 ncol()

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

7.4.3.20 nnozero()

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

7.4.3.21 nrow()

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

7.4.3.22 operator()() [1/2]

7.4.3.23 operator()() [2/2]

7.4.3.24 operator*=()

7.4.3.25 operator+=() [1/3]

7.4.3.26 operator+=() [2/3]

7.4.3.27 operator+=() [3/3]

7.4.3.28 operator-=() [1/3]

7.4.3.29 operator-=() [2/3]

7.4.3.30 operator-=() [3/3]

7.4.3.31 operator/=()

7.4.3.32 operator=() [1/2]

Move assignment.

7.4.3.33 operator=() [2/2]

Assignment constructor.

7.4.3.34 operator==()

7.4.3.35 out_of_range()

7.4.3.36 print()

7.4.3.37 reserve()

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

7.4.3.38 resize()

7.4.3.39 rm_cell()

7.4.3.40 row() [1/2]

7.4.3.41 row() [2/2]

7.4.3.42 rowsum()

7.4.3.43 set_data()

Set the data object.

Parameters

data_	
delete_ <i>←</i>	
data_	

7.4.3.44 swap_cells()

7.4.3.45 swap_cols()

7.4.3.46 swap_rows()

7.4.3.47 toggle_cell()

7.4.3.48 toggle_lock()

7.4.3.49 transpose()

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

7.4.3.50 zero_col()

7.4.3.51 zero_row()

7.4.4 Friends And Related Function Documentation

7.4.4.1 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.

7.4.4.2 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.

7.4.4.3 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.

7.4.4.4 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.

7.4.4.5 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.

7.4.5 Member Data Documentation

7.4.5.1 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 64 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

7.5 BArrayDenseCell< Cell_Type, Data_Type > Class Template Reference

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

Public Member Functions

- BArrayDenseCell (BArrayDense< Cell_Type, Data_Type > *Array_, uint i_, uint i_, bool check_bounds=true)
- ∼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 >

7.5.1 Detailed Description

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

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

7.5.2 Constructor & Destructor Documentation

7.5.2.1 BArrayDenseCell()

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

7.5.2.2 ~BArrayDenseCell()

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

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

7.5.3 Member Function Documentation

7.5.3.1 operator Cell_Type()

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

Definition at line 57 of file barraydensecell-meat.hpp.

7.5.3.2 operator*=()

Definition at line 37 of file barraydensecell-meat.hpp.

7.5.3.3 operator+=()

Definition at line 19 of file barraydensecell-meat.hpp.

7.5.3.4 operator-=()

Definition at line 28 of file barraydensecell-meat.hpp.

7.5.3.5 operator/=()

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

7.5.3.6 operator=()

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

7.5.3.7 operator==()

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

7.5.4 Friends And Related Function Documentation

7.5.4.1 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.

7.5.4.2 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.

7.5.4.3 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

7.6 BArrayDenseCell_const< Cell_Type, Data_Type > Class Template Reference

7.6.1 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

7.7 BArrayDenseCol< Cell_Type, Data_Type > Class Template Reference

#include <barraydensecol-bones.hpp>

Public Member Functions

- BArrayDenseCol (BArrayDense < Cell_Type, Data_Type > & array_, unsigned int j)
- Col_type< Cell_Type >::iterator & begin ()
- Col_type< Cell_Type >::iterator & end ()
- size_t size () const noexcept
- std::pair< unsigned int, Cell_Type * > & operator() (unsigned int i)

Friends

- class BArrayDense< Cell_Type, Data_Type >
- class BArrayDenseCell
 Cell Type, Data Type
- class BArrayDenseCell_const< Cell_Type, Data_Type >

7.7.1 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.

7.7.2 Constructor & Destructor Documentation

7.7.2.1 BArrayDenseCol()

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

7.7.3 Member Function Documentation

7.7.3.1 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.

7.7.3.2 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.

7.7.3.3 operator()()

```
template<typename Cell_Type = bool, typename Data_Type = bool> std::pair<unsigned int,Cell_Type*>& BArrayDenseCol< Cell_Type, Data_Type >::operator() ( unsigned int i ) [inline]
```

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

7.7.3.4 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.

7.7.4 Friends And Related Function Documentation

7.7.4.1 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.

7.7.4.2 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.

7.7.4.3 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-meat.hpp
- include/barry/barraydensecol-bones.hpp

7.8 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_, unsigned int j)
- Col_type< Cell_Type >::iterator begin ()
- Col_type< Cell_Type >::iterator end ()
- size t size () const noexcept
- const std::pair< unsigned int, Cell_Type * > operator() (unsigned int i) const

Friends

- class BArrayDenseCell
 Cell_Type, Data_Type
- class BArrayDenseCell_const< Cell_Type, Data_Type >

7.8.1 Detailed Description

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

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

7.8.2 Constructor & Destructor Documentation

7.8.2.1 BArrayDenseCol_const()

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

7.8.3 Member Function Documentation

7.8.3.1 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.

7.8.3.2 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.

7.8.3.3 operator()()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
const std::pair<unsigned int,Cell_Type*> BArrayDenseCol_const< Cell_Type, Data_Type >::operator()
(
    unsigned int i ) const [inline]
```

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

7.8.3.4 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.

7.8.4 Friends And Related Function Documentation

7.8.4.1 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.

7.8.4.2 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-meat.hpp
- include/barry/barraydensecol-bones.hpp

7.9 BArrayDenseRow< Cell_Type, Data_Type > Class Template Reference

#include <barraydenserow-bones.hpp>

Public Member Functions

- BArrayDenseRow (BArrayDense < Cell_Type, Data_Type > & array_, unsigned int i)
- Row_type< Cell_Type >::iterator & begin ()
- Row_type< Cell_Type >::iterator & end ()
- size_t size () const noexcept
- std::pair< unsigned int, Cell< Cell_Type > > & operator() (unsigned int i)

Friends

- class BArrayDense< Cell_Type, Data_Type >
- class BArrayDenseCell
 Cell_Type, Data_Type
- class BArrayDenseCell_const< Cell_Type, Data_Type >

7.9.1 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.

7.9.2 Constructor & Destructor Documentation

7.9.2.1 BArrayDenseRow()

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

7.9.3 Member Function Documentation

7.9.3.1 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.

7.9.3.2 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.

7.9.3.3 operator()()

```
\label{template} $$ \text{template}$ $$ \text{typename Cell_Type = bool, typename Data_Type = bool} $$ \text{std::pair}$ \text{cunsigned int,Cell}$ $$ \text{Cell_Type}$ > & $$ BArrayDenseRow$ Cell_Type, Data_Type >::operator() $$ ($$ unsigned int $i$ ) [inline] $$
```

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

7.9.3.4 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.

7.9.4 Friends And Related Function Documentation

7.9.4.1 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.

7.9.4.2 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.

7.9.4.3 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

7.10 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_, unsigned int 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< unsigned int, Cell< Cell_Type >> operator() (unsigned int i) const

Friends

- class BArrayDenseCell
 Cell_Type, Data_Type
- class BArrayDenseCell_const< Cell_Type, Data_Type >

7.10.1 Detailed Description

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

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

7.10.2 Constructor & Destructor Documentation

7.10.2.1 BArrayDenseRow_const()

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

7.10.3 Member Function Documentation

7.10.3.1 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.

7.10.3.2 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.

7.10.3.3 operator()()

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

7.10.3.4 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.

7.10.4 Friends And Related Function Documentation

7.10.4.1 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.

7.10.4.2 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

7.11 BArrayRow < Cell_Type, Data_Type > Class Template Reference

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

Public Member Functions

- BArrayRow (BArray< Cell_Type, Data_Type > *Array_, uint 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

7.11.1 Detailed Description

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

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

7.11.2 Constructor & Destructor Documentation

7.11.2.1 BArrayRow()

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

7.11.2.2 ∼BArrayRow()

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

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

7.11.3 Member Function Documentation

7.11.3.1 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
```

7.11.3.2 operator*=()

7.11.3.3 operator+=()

7.11.3.4 operator-=()

7.11.3.5 operator/=()

7.11.3.6 operator=()

7.11.3.7 operator==()

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

• include/barry/barrayrow-bones.hpp

7.12 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_, uint 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

7.12.1 Detailed Description

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

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

7.12.2 Constructor & Destructor Documentation

7.12.2.1 BArrayRow_const()

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

7.12.2.2 ~BArrayRow_const()

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

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

7.12.3 Member Function Documentation

7.12.3.1 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
```

7.12.3.2 operator"!=()

7.12.3.3 operator<()

7.12.3.4 operator<=()

7.12.3.5 operator==()

7.12.3.6 operator>()

7.12.3.7 operator>=()

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

• include/barry/barrayrow-bones.hpp

7.13 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_, uint &dim_ uint &i_, bool check_bounds=true)

 Construct a new BArrayVector object.
- ∼BArrayVector ()
- bool is_row () const noexcept
- bool is_col () const noexcept
- uint 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

7.13.1 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 13 of file barrayvector-bones.hpp.

7.13.2 Constructor & Destructor Documentation

7.13.2.1 BArrayVector()

Construct a new BArrayVector object.

Parameters

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

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

7.13.2.2 ~BArrayVector()

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

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

7.13.3 Member Function Documentation

7.13.3.1 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 52 of file barrayvector-meat.hpp.

7.13.3.2 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 66 of file barrayvector-meat.hpp.

7.13.3.3 is_col()

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

Definition at line 36 of file barrayvector-meat.hpp.

7.13.3.4 is_row()

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

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

7.13.3.5 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 177 of file barrayvector-meat.hpp.

7.13.3.6 operator*=()

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

7.13.3.7 operator+=()

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

7.13.3.8 operator-=()

Definition at line 114 of file barrayvector-meat.hpp.

7.13.3.9 operator/=()

Definition at line 156 of file barrayvector-meat.hpp.

7.13.3.10 operator=()

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

7.13.3.11 operator==()

Definition at line 187 of file barrayvector-meat.hpp.

7.13.3.12 size()

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

Definition at line 41 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

7.14 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_, uint &dim_ uint &i_, bool check_bounds=true)
- ~BArrayVector_const ()
- bool is_row () const noexcept
- bool is_col () const noexcept
- uint 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

7.14.1 Detailed Description

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

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

7.14.2 Constructor & Destructor Documentation

7.14.2.1 BArrayVector_const()

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

7.14.2.2 ~BArrayVector_const()

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

Definition at line 110 of file barrayvector-bones.hpp.

7.14.3 Member Function Documentation

7.14.3.1 begin()

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

7.14.3.2 end()

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

7.14.3.3 is_col()

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

7.14.3.4 is_row()

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

7.14.3.5 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 214 of file barrayvector-meat.hpp.

7.14.3.6 operator"!=()

Definition at line 251 of file barrayvector-meat.hpp.

7.14.3.7 operator<()

Definition at line 256 of file barrayvector-meat.hpp.

7.14.3.8 operator<=()

Definition at line 283 of file barrayvector-meat.hpp.

7.14.3.9 operator==()

Definition at line 224 of file barrayvector-meat.hpp.

7.14.3.10 operator>()

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

7.14.3.11 operator>=()

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

7.14.3.12 size()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
uint 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

7.15 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 (unsigned int x)
- void add (int x)
- Cell ()
- Cell ()
- Cell ()

Public Attributes

- Cell_Type value
- bool visited
- · bool active

7.15.1 Detailed Description

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

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

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

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

7.15.2 Constructor & Destructor Documentation

7.15.2.1 Cell() [1/7]

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

7.15.2.2 Cell() [2/7]

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

7.15.2.3 ∼Cell()

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

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

7.15.2.4 Cell() [3/7]

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

7.15.2.5 Cell() [4/7]

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

7.15.2.6 Cell() [5/7]

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

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

7.15.2.7 Cell() [6/7]

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

Definition at line 65 of file cell-meat.hpp.

7.15.2.8 Cell() [7/7]

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

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

7.15.3 Member Function Documentation

7.15.3.1 add() [1/4]

7.15.3.2 add() [2/4]

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

7.15.3.3 add() [3/4]

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

Definition at line 54 of file cell-meat.hpp.

7.15.3.4 add() [4/4]

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

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

7.15.3.5 operator Cell_Type()

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

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

7.15.3.6 operator"!=()

Definition at line 33 of file cell-meat.hpp.

7.15.3.7 operator=() [1/2]

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

7.15.3.8 operator=() [2/2]

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

7.15.3.9 operator==()

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

7.15.4 Member Data Documentation

7.15.4.1 active

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

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

7.15.4.2 value

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

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

7.15.4.3 visited

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

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

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

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

7.16 Cell_const < Cell_Type > Class Template Reference

7.16.1 Detailed Description

```
\label{template} \begin{tabular}{ll} template < typename Cell\_Type > \\ class Cell\_const < Cell\_Type > \\ \end{tabular}
```

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

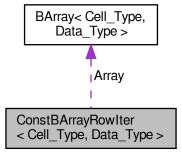
The documentation for this class was generated from the following file:

• include/barry/barray-meat.hpp

7.17 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

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

7.17.1 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.

7.17.2 Constructor & Destructor Documentation

7.17.2.1 ConstBArrayRowIter()

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

7.17.2.2 ~ConstBArrayRowlter()

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

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

7.17.3 Member Data Documentation

7.17.3.1 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.

7.17.3.2 current_col

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

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

7.17.3.3 current_row

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

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

7.17.3.4 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

7.18 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, uint i, uint j)
- double init (Array_Type &Array, uint i, uint 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 (Counter_fun_type< Array_Type, Data_Type > count_fun_, Counter_fun_type< Array_Type, Data_Type > init_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.

Public Attributes

- Counter_fun_type
 Array_Type, Data_Type > count_fun
- Counter_fun_type
 Array_Type, Data_Type > init_fun
- Data_Type data
- std::string name = ""
- std::string desc = ""

7.18.1 Detailed Description

```
\label{template} $$ \ensuremath{\sf template}$$ < typename \ensuremath{\sf Array\_Type}$ = BArray<>, typename \ensuremath{\sf Data\_Type}$ = bool> class \ensuremath{\sf Counter}< Array\_Type, \ensuremath{\sf Data\_Type}>
```

A counter function based on change statistics.

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

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

7.18.2 Constructor & Destructor Documentation

7.18.2.1 Counter() [1/4]

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

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

7.18.2.2 Counter() [2/4]

Definition at line 60 of file counters-bones.hpp.

7.18.2.3 Counter() [3/4]

Copy constructor.

7.18.2.4 Counter() [4/4]

Move constructor.

7.18.2.5 ∼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.

7.18.3 Member Function Documentation

7.18.3.1 count()

7.18.3.2 get_description()

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

7.18.3.3 get_name()

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

7.18.3.4 init()

7.18.3.5 operator=() [1/2]

Copy assignment.

7.18.3.6 operator=() [2/2]

Move assignment.

7.18.4 Member Data Documentation

7.18.4.1 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 41 of file counters-bones.hpp.

7.18.4.2 data

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

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

7.18.4.3 desc

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

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

7.18.4.4 init_fun

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

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

7.18.4.5 name

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

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

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

• include/barry/counters-bones.hpp

7.19 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[] (uint idx)

Returns a pointer to a particular counter.

• std::size_t size () const noexcept

Number of counters in the set.

- void add counter (Counter< Array Type, Data Type > counter)
- void add_counter (Counter_fun_type< Array_Type, Data_Type > count_fun_, Counter_fun_type< Array_
 —
 Type, Data_Type > init_fun_, Data_Type data_, std::string name_="", std::string desc_="")
- std::vector< std::string > get_names () const
- std::vector< std::string > get_descriptions () const

7.19.1 Detailed Description

```
\label{template} $$ \ensuremath{\sf template}$$ < 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 95 of file counters-bones.hpp.

7.19.2 Constructor & Destructor Documentation

7.19.2.1 Counters() [1/3]

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

7.19.2.2 ∼Counters()

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

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

7.19.2.3 Counters() [2/3]

Copy constructor.

Parameters



7.19.2.4 Counters() [3/3]

Move constructor.

Parameters

counters⇔

7.19.3 Member Function Documentation

7.19.3.1 add_counter() [1/2]

7.19.3.2 add_counter() [2/2]

7.19.3.3 get_descriptions()

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

7.19.3.4 get_names()

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

7.19.3.5 operator=() [1/2]

Copy assignment constructor.

Parameters

counter←	

Returns

Counters<Array_Type,Data_Type>

7.19.3.6 operator=() [2/2]

Move assignment constructor.

Parameters



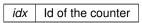
Returns

Counters<Array_Type,Data_Type>&

7.19.3.7 operator[]()

Returns a pointer to a particular counter.

Parameters



Returns

Counter<Array_Type,Data_Type>*

7.19.3.8 size()

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

Number of counters in the set.

Returns

uint

Definition at line 150 of file counters-bones.hpp.

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

• include/barry/counters-bones.hpp

7.20 DEFMCounterData Class Reference

Data class used to store arbitrary uint or double vectors.

```
#include <defm.hpp>
```

Public Member Functions

- DEFMCounterData ()
- DEFMCounterData (const std::vector< uint > indices_, const std::vector< double > numbers_, size_←
 t markov_order_)
- ∼DEFMCounterData ()

Public Attributes

- std::vector< size_t > indices
- std::vector< double > numbers
- size_t markov_order

Order of the markov process.

7.20.1 Detailed Description

Data class used to store arbitrary uint or double vectors.

Definition at line 73 of file defm.hpp.

7.20.2 Constructor & Destructor Documentation

7.20.2.1 **DEFMCounterData()** [1/2]

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

Definition at line 80 of file defm.hpp.

7.20.2.2 **DEFMCounterData()** [2/2]

```
DEFMCounterData::DEFMCounterData (
    const std::vector< uint > indices_,
    const std::vector< double > numbers_,
    size_t markov_order_ ) [inline]
```

Definition at line 81 of file defm.hpp.

7.20.2.3 ~DEFMCounterData()

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

Definition at line 87 of file defm.hpp.

7.20.3 Member Data Documentation

7.20.3.1 indices

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

Definition at line 76 of file defm.hpp.

7.20.3.2 markov_order

```
size_t DEFMCounterData::markov_order
```

Order of the markov process.

Definition at line 78 of file defm.hpp.

7.20.3.3 numbers

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

Definition at line 77 of file defm.hpp.

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

• include/barry/counters/defm.hpp

7.21 DEFMData Class Reference

Data class for DEFM arrays.

```
#include <defm.hpp>
```

Public Member Functions

```
• DEFMData ()
```

 DEFMData (std::vector< double > *covariates_, size_t obs_start_, size_t obs_n_times_, size_t n_states_, size_t n_covariates_)

Constructor.

∼DEFMData ()

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

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

double at (size_t i, size_t j)

Public Attributes

std::vector< double > * covariates

Vector of covariates (complete vector)

· size_t obs_start

Index of the observation in the data.

size_t obs_n_times

Number of observations in the model.

size_t n_covariates

Number of covariates included in the model.

7.21.1 Detailed Description

Data class for DEFM arrays.

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

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 20 of file defm.hpp.

7.21.2 Constructor & Destructor Documentation

7.21.2.1 **DEFMData()** [1/2]

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

Definition at line 28 of file defm.hpp.

7.21.2.2 DEFMData() [2/2]

Constructor.

Parameters

covariates_	Pointer to the attribute data.
obs_start_	Location of the current observation in the covariates vector
obs_n_←	Number of observations in the model
times_	
<i>n</i> _⇔	Number of columns (covariates.)
covariates_	

Definition at line 38 of file defm.hpp.

7.21.2.3 \sim DEFMData()

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

Definition at line 57 of file defm.hpp.

7.21.3 Member Function Documentation

7.21.3.1 at()

Definition at line 66 of file defm.hpp.

7.21.3.2 operator()()

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

Parameters



Returns

double

Definition at line 61 of file defm.hpp.

7.21.4 Member Data Documentation

7.21.4.1 covariates

```
std::vector< double >* DEFMData::covariates
```

Vector of covariates (complete vector)

Definition at line 23 of file defm.hpp.

7.21.4.2 n_covariates

```
size_t DEFMData::n_covariates
```

Number of covariates included in the model.

Definition at line 26 of file defm.hpp.

7.21.4.3 obs_n_times

```
size_t DEFMData::obs_n_times
```

Number of observations in the model.

Definition at line 25 of file defm.hpp.

7.21.4.4 obs_start

```
size_t DEFMData::obs_start
```

Index of the observation in the data.

Definition at line 24 of file defm.hpp.

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

· include/barry/counters/defm.hpp

7.22 Entries < Cell_Type > Class Template Reference

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

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

Public Member Functions

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

Public Attributes

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

7.22.1 Detailed Description

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

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

Template Parameters

```
Cell_Type Any type
```

Definition at line 79 of file typedefs.hpp.

7.22.2 Constructor & Destructor Documentation

7.22.2.1 Entries() [1/2]

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

Definition at line 85 of file typedefs.hpp.

7.22.2.2 Entries() [2/2]

Definition at line 86 of file typedefs.hpp.

7.22.2.3 ∼Entries()

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

Definition at line 93 of file typedefs.hpp.

7.22.3 Member Function Documentation

7.22.3.1 resize()

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

Definition at line 95 of file typedefs.hpp.

7.23 Flock Class Reference 117

7.22.4 Member Data Documentation

7.22.4.1 source

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

Definition at line 81 of file typedefs.hpp.

7.22.4.2 target

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

Definition at line 82 of file typedefs.hpp.

7.22.4.3 val

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

Definition at line 83 of file typedefs.hpp.

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

• include/barry/typedefs.hpp

7.23 Flock Class Reference

A Flock is a group of Geese.

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

Public Member Functions

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

Add a tree to the flock.

· void set seed (const unsigned int &s)

Set the seed of the model.

- void init (unsigned int bar width=BARRY PROGRESS BAR WIDTH)
- phylocounters::PhyloCounters * get_counters ()
- phylocounters::PhyloSupport * get support fun ()
- std::vector< std::vector< double > > * get_stats_support ()
- std::vector< std::vector< double > > * get stats target ()
- phylocounters::PhyloModel * get model ()

Returns the joint likelihood of the model.

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

Access the i-th geese element.

Information about the model

- · unsigned int nfuns () const noexcept
- unsigned int ntrees () const noexcept
- std::vector< unsigned int > nnodes () const noexcept
- std::vector< unsigned int > nleafs () const noexcept
- unsigned int nterms () const
- unsigned int support_size () const noexcept
- std::vector< std::string > colnames () const
- unsigned int 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
- unsigned int nfunctions = 0u
- bool initialized = false
- std::mt19937 rengine
- phylocounters::PhyloModel model = phylocounters::PhyloModel()

7.23.1 Detailed Description

A Flock is a group of Geese.

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

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

7.23 Flock Class Reference 119

7.23.2 Constructor & Destructor Documentation

7.23.2.1 Flock()

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

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

7.23.2.2 ∼Flock()

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

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

7.23.3 Member Function Documentation

7.23.3.1 add_data()

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

Add a tree to the flock.

Parameters

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

Returns

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

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

7.23.3.2 colnames()

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

Definition at line 224 of file flock-meat.hpp.

7.23.3.3 get_counters()

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

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

7.23.3.4 get_model()

```
phylocounters::PhyloModel * Flock::get_model ( ) [inline]
```

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

7.23.3.5 get_stats_support()

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

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

7.23.3.6 get_stats_target()

```
std::vector< std::vector< double > > * Flock::get_stats_target ( ) [inline]
```

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

7.23.3.7 get_support_fun()

```
phylocounters::PhyloSupport * Flock::get_support_fun ( ) [inline]
```

Definition at line 110 of file flock-meat.hpp.

7.23 Flock Class Reference 121

7.23.3.8 init()

```
void Flock::init (
     unsigned int bar_width = BARRY_PROGRESS_BAR_WIDTH ) [inline]
```

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

7.23.3.9 likelihood_joint()

Returns the joint likelihood of the model.

Parameters

par	Vector of model parameters.	
as_log	When true it will return the value as log.	
use_reduced_sequence	, , ,	
	which is faster.	

Returns

double

Definition at line 138 of file flock-meat.hpp.

7.23.3.10 nfuns()

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

Definition at line 167 of file flock-meat.hpp.

7.23.3.11 nleafs()

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

Definition at line 195 of file flock-meat.hpp.

7.23.3.12 nnodes()

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

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

7.23.3.13 nterms()

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

Definition at line 209 of file flock-meat.hpp.

7.23.3.14 ntrees()

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

Definition at line 174 of file flock-meat.hpp.

7.23.3.15 operator()()

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

Access the i-th geese element.

Parameters

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

Returns

Geese*

Definition at line 302 of file flock-meat.hpp.

7.23.3.16 parse_polytomies()

```
unsigned int Flock::parse_polytomies (
          bool verb = true,
          std::vector< size_t > * dist = nullptr ) const [inline], [noexcept]
```

7.23 Flock Class Reference 123

Check polytomies and return the largest.

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

7.23.3.17 print()

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

Definition at line 258 of file flock-meat.hpp.

7.23.3.18 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.

7.23.3.19 support_size()

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

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

7.23.4 Member Data Documentation

7.23.4.1 dat

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

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

7.23.4.2 initialized

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

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

7.23.4.3 model

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

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

7.23.4.4 nfunctions

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

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

7.23.4.5 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

7.24 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< double > &x) const
```

7.24.1 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 27 of file freqtable.hpp.

7.24.2 Constructor & Destructor Documentation

7.24.2.1 FreqTable()

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

Definition at line 39 of file freqtable.hpp.

7.24.2.2 ∼FreqTable()

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

Definition at line 40 of file freqtable.hpp.

7.24.3 Member Function Documentation

7.24.3.1 add()

Definition at line 64 of file freqtable.hpp.

7.24.3.2 as_vector()

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

Definition at line 144 of file freqtable.hpp.

7.24.3.3 clear()

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

Definition at line 173 of file freqtable.hpp.

7.24.3.4 get_data()

```
template<typename T = double>
const std::vector< double >& FreqTable< T >::get_data ( ) const [inline]
```

Definition at line 45 of file freqtable.hpp.

7.24.3.5 get_index()

```
template<typename T = double>
const std::unordered_map<size_t,size_t>& FreqTable< T >::get_index ( ) const [inline]
```

Definition at line 46 of file freqtable.hpp.

7.24.3.6 make_hash()

Definition at line 244 of file freqtable.hpp.

7.24.3.7 print()

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

Definition at line 209 of file freqtable.hpp.

7.24.3.8 reserve()

Definition at line 187 of file freqtable.hpp.

7.24.3.9 size()

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

Number of unique elements in the table. (.

Returns

size_t

Definition at line 236 of file freqtable.hpp.

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

include/barry/freqtable.hpp

7.25 Geese Class Reference

Annotated Phylo Model.

#include <defm-bones.hpp>

Public Member Functions

- ∼Geese ()
- void init (unsigned int bar width=BARRY PROGRESS BAR WIDTH)
- void inherit_support (const Geese &model_, bool delete_support_=false)
- void calc sequence (Node *n=nullptr)
- void calc reduced sequence ()
- double likelihood (const std::vector< double > &par, bool as_log=false, bool use_reduced_sequence=true)
- double likelihood_exhaust (const std::vector< double > &par)
- std::vector< double > get_probabilities () const
- void set_seed (const unsigned int &s)
- std::vector< std::vector< unsigned int > > simulate (const std::vector< double > &par)
- std::vector< std::vector< double > > observed counts ()
- void print_observed_counts ()
- void print () const

Prints information about the DEFM.

- void init node (Node &n)
- void update_annotations (unsigned int nodeid, std::vector< unsigned int > newann)
- std::vector< std::vector< bool >> get_states () const

Powerset of a gene's possible states.

• std::vector< unsigned int > get_annotated_nodes () const

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

- ∼Geese ()
- · void init (unsigned int bar width=BARRY PROGRESS BAR WIDTH)
- void inherit_support (const Geese &model_, bool delete_support_=false)
- void calc_sequence (Node *n=nullptr)
- void calc reduced sequence ()
- double likelihood (const std::vector< double > &par, bool as_log=false, bool use_reduced_sequence=true)
- double likelihood_exhaust (const std::vector< double > &par)
- std::vector< double > get_probabilities () const
- void set_seed (const unsigned int &s)
- std::vector< std::vector< unsigned int > > simulate (const std::vector< double > &par)
- std::vector< std::vector< double >> observed counts ()
- void print_observed_counts ()
- · void print () const

Prints information about the GEESE.

- void init_node (Node &n)
- void update_annotations (unsigned int nodeid, std::vector< unsigned int > newann)
- std::vector< std::vector< bool > > get states () const

Powerset of a gene's possible states.

• std::vector< unsigned int > get_annotated_nodes () const

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

Construct a new Geese object

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

Parameters

annotations	A vector of vectors with annotations. It should be of length k (number of functions). Each vector should be of length N (equal to the number of nodes, including interior). Possible values are 0, 1, and 9.
geneid	Id of the gene. It should be of length ${ m N}.$
parent	Id of the parent gene. Also of length ${\tt N}$
duplication	Logical scalar indicating the type of event (true: duplication, false: speciation.)

The ordering of the entries does not matter. Passing the nodes in post order or not makes no difference to the constructor.

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

Information about the model

Parameters

verb When true it will print out information about the encountered polytomies.

· unsigned int nfuns () const noexcept

Number of functions analyzed.

· unsigned int nnodes () const noexcept

Number of nodes (interior + leaf)

· unsigned int nleafs () const noexcept

Number of leaf.

• unsigned int nterms () const

Number of terms included.

• unsigned int support_size () const noexcept

Number of unique sets of sufficient stats.

std::vector< unsigned int > nannotations () const noexcept

Number of annotations.

std::vector< std::string > colnames () const

Names of the terms in the model.

unsigned int parse_polytomies (bool verb=true, std::vector < size_t > *dist=nullptr) const noexcept
 Check polytomies and return the largest.

· unsigned int nfuns () const noexcept

Number of functions analyzed.

• unsigned int nnodes () const noexcept

Number of nodes (interior + leaf)

unsigned int nleafs () const noexcept

Number of leaf.

unsigned int nterms () const

Number of terms included.

unsigned int support_size () const noexcept

Number of unique sets of sufficient stats.

std::vector< unsigned int > nannotations () const noexcept

Number of annotations.

std::vector< std::string > colnames () const

Names of the terms in the model.

unsigned int 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_backend (const std::vector < double > &par, bool use_← reduced_sequence, const std::vector < uint > &preorder)
- std::vector< std::vector< double > > predict_exhaust_backend (const std::vector< double > &par, const std::vector< uint > &preorder)
- std::vector< std::vector< double > > predict_exhaust (const std::vector< double > &par)
- std::vector< std::vector< double > > predict_sim (const std::vector< double > &par, bool only_← annotated=false, unsigned int nsims=10000u)
- 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< uint > &preorder)
- std::vector< std::vector< double >> predict_exhaust_backend (const std::vector< double > &par, const std::vector< uint > &preorder)
- std::vector< std::vector< double > > predict_exhaust (const std::vector< double > &par)
- std::vector < std::vector < double > > predict_sim (const std::vector < double > &par, bool only_
 —
 annotated=false, unsigned int 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::PhyloCounters * get_counters()
phylocounters::PhyloSupport * get_support_fun()
std::mt19937 * get_rengine()
phylocounters::PhyloCounters * get_counters()
phylocounters::PhyloCounters * get_support_fun()
phylocounters::PhyloModel * get_model()
phylocounters::PhyloSupport * get_support_fun()
```

Public Attributes

- · unsigned int nfunctions
- std::map< unsigned int, Node > nodes
- barry::MapVec type< unsigned int > map to nodes
- $std::vector < std::vector < std::vector < size_t >> pset_loc$

Locations of columns.

- std::vector< unsigned int > sequence
- std::vector< unsigned int > reduced_sequence
- bool initialized = false
- bool delete_rengine = false
- bool delete_support = false

7.25.1 Detailed Description

Annotated Phylo Model.

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

Definition at line 65 of file defm-bones.hpp.

7.25.2 Constructor & Destructor Documentation

7.25.2.1 Geese() [1/8]

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

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

7.25.2.2 Geese() [2/8]

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

Definition at line 20 of file geese-meat-constructors.hpp.

7.25.2.3 Geese() [3/8]

Definition at line 214 of file geese-meat-constructors.hpp.

7.25.2.4 Geese() [4/8]

Definition at line 293 of file geese-meat-constructors.hpp.

7.25.2.5 ~Geese() [1/2]

```
Geese::∼Geese ( ) [inline]
```

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

7.25.2.6 Geese() [5/8]

```
Geese::Geese ( )
```

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7.25.2.7 Geese() [6/8]

```
Geese::Geese (
         std::vector< std::vector< unsigned int > > & annotations,
         std::vector< unsigned int > & geneid,
         std::vector< int > & parent,
         std::vector< bool > & duplication )
```

7.25.2.8 Geese() [7/8]

7.25.2.9 Geese() [8/8]

7.25.2.10 ~Geese() [2/2]

```
Geese::∼Geese ( )
```

7.25.3 Member Function Documentation

7.25.3.1 calc_reduced_sequence() [1/2]

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

Definition at line 331 of file defm-meat.hpp.

7.25.3.2 calc_reduced_sequence() [2/2]

```
void Geese::calc_reduced_sequence ( )
```

7.25.3.3 calc_sequence() [1/2]

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

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

7.25.3.4 calc_sequence() [2/2]

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

7.25.3.5 colnames() [1/2]

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

Names of the terms in the model.

Definition at line 453 of file defm-meat.hpp.

7.25.3.6 colnames() [2/2]

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

Names of the terms in the model.

7.25.3.7 get annotated nodes() [1/2]

```
std::vector< unsigned int > Geese::get_annotated_nodes ( ) const [inline]
```

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

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

7.25.3.8 get_annotated_nodes() [2/2]

```
{\tt std::vector} < {\tt unsigned int} > {\tt Geese::get\_annotated\_nodes} ( ) const
```

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

7.25 Geese Class Reference 135

7.25.3.9 get_counters() [1/2]

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

Definition at line 652 of file defm-meat.hpp.

7.25.3.10 get_counters() [2/2]

```
phylocounters::PhyloCounters* Geese::get_counters ( )
```

7.25.3.11 get_model() [1/2]

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

Definition at line 657 of file defm-meat.hpp.

7.25.3.12 get_model() [2/2]

```
phylocounters::PhyloModel* Geese::get_model ( )
```

7.25.3.13 get_probabilities() [1/2]

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

Definition at line 379 of file defm-meat.hpp.

7.25.3.14 get_probabilities() [2/2]

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

7.25.3.15 get_rengine() [1/2]

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

Definition at line 647 of file defm-meat.hpp.

7.25.3.16 get_rengine() [2/2]

```
std::mt19937* Geese::get_rengine ( )
```

7.25.3.17 get_states() [1/2]

```
std::vector< std::vector< bool > > Geese::get_states ( ) const [inline]
```

Powerset of a gene's possible states.

This list of vectors is used throughout Geese. It lists all possible combinations of functional states for any gene. Thus, for P functions, there will be 2^{P} possible combinations.

Returns

```
std::vector< std::vector< bool >> of length 2^{^{\text{}}}P.
```

Definition at line 665 of file defm-meat.hpp.

7.25.3.18 get_states() [2/2]

```
std::vector< std::vector< bool > > Geese::get_states ( ) const
```

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.
```

7.25.3.19 get_support_fun() [1/2]

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

Definition at line 661 of file defm-meat.hpp.

7.25.3.20 get_support_fun() [2/2]

```
phylocounters::PhyloSupport* Geese::get_support_fun ( )
```

7.25 Geese Class Reference 137

7.25.3.21 inherit_support() [1/2]

Definition at line 230 of file defm-meat.hpp.

7.25.3.22 inherit_support() [2/2]

7.25.3.23 init() [1/2]

```
void Geese::init (
          unsigned int bar_width = BARRY_PROGRESS_BAR_WIDTH ) [inline]
```

Definition at line 103 of file defm-meat.hpp.

7.25.3.24 init() [2/2]

```
void Geese::init (
     unsigned int bar_width = BARRY_PROGRESS_BAR_WIDTH )
```

7.25.3.25 init_node() [1/2]

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

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

7.25.3.26 init_node() [2/2]

```
void Geese::init_node (
    Node & n )
```

7.25.3.27 likelihood() [1/2]

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

7.25.3.28 likelihood() [2/2]

7.25.3.29 likelihood exhaust() [1/2]

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

7.25.3.30 likelihood_exhaust() [2/2]

7.25.3.31 nannotations() [1/2]

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

Number of annotations.

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

7.25 Geese Class Reference 139

7.25.3.32 nannotations() [2/2]

```
std::vector< unsigned int > Geese::nannotations ( ) const [noexcept]
```

Number of annotations.

7.25.3.33 nfuns() [1/2]

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

Number of functions analyzed.

Definition at line 400 of file defm-meat.hpp.

7.25.3.34 nfuns() [2/2]

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

Number of functions analyzed.

7.25.3.35 nleafs() [1/2]

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

Number of leaf.

Definition at line 414 of file defm-meat.hpp.

7.25.3.36 nleafs() [2/2]

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

Number of leaf.

7.25.3.37 nnodes() [1/2]

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

Number of nodes (interior + leaf)

Definition at line 407 of file defm-meat.hpp.

7.25.3.38 nnodes() [2/2]

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

Number of nodes (interior + leaf)

7.25.3.39 nterms() [1/2]

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

Number of terms included.

Definition at line 426 of file defm-meat.hpp.

7.25.3.40 nterms() [2/2]

```
unsigned int Geese::nterms ( ) const
```

Number of terms included.

7.25.3.41 observed_counts() [1/2]

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

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

7.25.3.42 observed_counts() [2/2]

```
std::vector< std::vector<double> > Geese::observed_counts ( )
```

7.25.3.43 operator=() [1/4]

7.25 Geese Class Reference 141

7.25.3.44 operator=() [2/4]

7.25.3.45 operator=() [3/4]

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

7.25.3.46 operator=() [4/4]

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

7.25.3.47 parse_polytomies() [1/2]

```
unsigned int Geese::parse_polytomies (
          bool verb = true,
          std::vector< size_t > * dist = nullptr ) const [inline], [noexcept]
```

Check polytomies and return the largest.

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

7.25.3.48 parse_polytomies() [2/2]

Check polytomies and return the largest.

7.25.3.49 predict() [1/2]

Definition at line 243 of file geese-meat-predict.hpp.

7.25.3.50 predict() [2/2]

7.25.3.51 predict_backend() [1/2]

< True if the array belongs to the set

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

7.25.3.52 predict_backend() [2/2]

7.25.3.53 predict_exhaust() [1/2]

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

7.25.3.54 predict_exhaust() [2/2]

7.25 Geese Class Reference 143

7.25.3.55 predict_exhaust_backend() [1/2]

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

7.25.3.56 predict_exhaust_backend() [2/2]

7.25.3.57 predict_sim() [1/2]

```
std::vector< std::vector< double > > Geese::predict_sim (
    const std::vector< double > & par,
    bool only_annotated = false,
    unsigned int nsims = 10000u ) [inline]
```

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

7.25.3.58 predict_sim() [2/2]

7.25.3.59 print() [1/2]

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

Prints information about the DEFM.

Definition at line 629 of file defm-meat.hpp.

7.25.3.60 print() [2/2]

```
void Geese::print ( ) const
```

Prints information about the GEESE.

7.25.3.61 print_observed_counts() [1/2]

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

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

7.25.3.62 print_observed_counts() [2/2]

```
void Geese::print_observed_counts ( )
```

7.25.3.63 set_seed() [1/2]

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

7.25.3.64 set_seed() [2/2]

7.25.3.65 simulate() [1/2]

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

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

7.25 Geese Class Reference 145

7.25.3.66 simulate() [2/2]

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

7.25.3.67 support_size() [1/2]

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

Number of unique sets of sufficient stats.

Definition at line 434 of file defm-meat.hpp.

7.25.3.68 support_size() [2/2]

```
unsigned int Geese::support_size ( ) const [noexcept]
```

Number of unique sets of sufficient stats.

7.25.3.69 update_annotations() [1/2]

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

Definition at line 258 of file defm-meat.hpp.

7.25.3.70 update_annotations() [2/2]

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

7.25.4 Member Data Documentation

7.25.4.1 delete_rengine

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

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

7.25.4.2 delete_support

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

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

7.25.4.3 initialized

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

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

7.25.4.4 map_to_nodes

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

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

7.25.4.5 nfunctions

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

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

7.25.4.6 nodes

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

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

7.25.4.7 pset_loc

std::vector< std::vector< size_t > > Seese::pset_loc

Locations of columns.

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

7.25.4.8 reduced_sequence

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

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

7.25.4.9 sequence

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

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

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

- include/barry/models/defm/defm-bones.hpp
- include/barry/models/geese/geese-bones.hpp
- include/barry/models/defm/defm-meat.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

7.26 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>

Public Member Functions

- void set_rengine (std::mt19937 *rengine_, bool delete_=false)
- void set_seed (unsigned int s)
- Model ()
- Model (uint 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_)
- ∼Model ()
- · void store psets () noexcept
- void set_keygen (std::function < std::vector < double >(const Array_Type &) > keygen_)
- std::vector< double > gen_key (const Array_Type &Array_)
- uint add_array (const Array_Type &Array_, bool force_new=false)

Adds an array to the support of not already included.

- · void print stats (uint i) const
- · void print () const

Prints information about the model.

- Array_Type sample (const Array_Type &Array_, const std::vector< double > ¶ms={})
- Array Type sample (const uint &i, const std::vector< double > ¶ms)
- double conditional_prob (const Array_Type &Array_, const std::vector< double > ¶ms, unsigned int i, unsigned int 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_)

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

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

- void add_rule (Rule < Array_Type, Data_Rule_Type > &rule)
- void add_rule (Rule_fun_type< Array_Type, Data_Rule_Type > count_fun_, Data_Rule_Type data_)
- void set_rules (Rules< Array_Type, Data_Rule_Type > *rules_)
- void add_rule_dyn (Rule< Array_Type, Data_Rule_Dyn_Type > &rule)
- void 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 uint &i, bool as_log=false)
- double likelihood (const std::vector < double > &target_, const uint &i, bool as log=false)
- double likelihood (const std::vector < double > ¶ms, const double *target_, const uint &i, bool as_←
 log=false)
- double likelihood total (const std::vector< double > ¶ms, bool as log=false)

Extract elements by index

Parameters

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

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

- unsigned int size () const noexcept
- unsigned int size unique () const noexcept
- · unsigned int nterms () const noexcept
- unsigned int support_size () const noexcept
- std::vector< std::string > colnames () const
- std::vector< std::vector< double > > * get_stats_target ()

Raw pointers to the support and target statistics.

- std::vector< std::vector< double >> * get_stats_support ()
- std::vector< unsigned int > * get_arrays2support ()
- std::vector< std::vector< Array_Type > > * get_pset_arrays ()
- std::vector< std::vector< double > > * get_pset_stats ()

Statistics of the support(s)

std::vector< std::vector< double >> * get_pset_probs ()

void set_transform_model (std::function< std::vector< double >(double *, unsigned int)> fun, std::vector< std::string > names)

Set the transform_model_fun object.

std::vector< double > transform model (double *data, unsigned int k)

7.26.1 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:

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

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

Template Parameters

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

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

7.26.2 Constructor & Destructor Documentation

7.26.2.1 Model() [1/3]

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::Model ()
```

7.26.2.2 Model() [2/3]

7.26.2.3 Model() [3/3]

7.26.2.4 ~Model()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>

Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::~Model ( ) [inline]
```

Definition at line 180 of file model-bones.hpp.

7.26.3 Member Function Documentation

7.26.3.1 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.

7.26.3.2 add_counter() [1/2]

7.26.3.3 add_counter() [2/2]

7.26.3.4 add_rule() [1/2]

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::add_rule (
Rule< Array_Type, Data_Rule_Type > & rule )
```

7.26.3.5 add rule() [2/2]

7.26.3.6 add_rule_dyn() [1/2]

7.26.3.7 add_rule_dyn() [2/2]

7.26.3.8 colnames()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::string > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_\times
Dyn_Type >::colnames ( ) const
```

7.26.3.9 conditional_prob()

Conditional probability ("Gibbs sampler")

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

7.26.3.10 gen_key()

7.26.3.11 get_arrays2support()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< unsigned int >* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_←
Rule_Dyn_Type >::get_arrays2support ()
```

7.26.3.12 get_counters()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Counters<Array_Type, Data_Counter_Type>* Model< Array_Type, Data_Counter_Type, Data_Rule_Dyn_Type >::get_counters ( )
```

7.26.3.13 get_norm_const()

7.26.3.14 get pset()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarrow Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> const std::vector< Array_Type >* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data\leftarrow _Rule_Dyn_Type >::get_pset ( const uint & i )
```

7.26.3.15 get_pset_arrays()

```
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< Array_Type > >* Model< Array_Type, Data_Counter_Type, Data_Rule_\times
Type, Data_Rule_Dyn_Type >::get_pset_arrays ()
```

7.26.3.16 get_pset_probs()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector<double> >* Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_pset_probs ()
```

7.26.3.17 get_pset_stats() [1/2]

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector<double> >* Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_pset_stats ()
```

Statistics of the support(s)

7.26.3.18 get pset stats() [2/2]

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarrow Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> const std::vector< double >* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_\leftarrow Rule_Dyn_Type >::get_pset_stats ( const uint & i )
```

7.26.3.19 get_rengine()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
const std::mt19937* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type
>::get_rengine ( ) const
```

7.26.3.20 get_rules()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Rules<Array_Type, Data_Rule_Type>* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data←
_Rule_Dyn_Type >::get_rules ()
```

7.26.3.21 get_rules_dyn()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Rules<Array_Type, Data_Rule_Dyn_Type>* Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_rules_dyn ()
```

7.26.3.22 get_stats_support()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector< double > >* Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_stats_support ()
```

7.26.3.23 get_stats_target()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector< double > >* Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_stats_target ()
```

Raw pointers to the support and target statistics.

The support of the model is stored as a vector of vector < double>. Each element of it contains the support for an specific type of array included. It represents an array of size $(k + 1) \times n$ 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.

7.26.3.24 get support fun()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Support<Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type>* Model< Array_Type,
Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::get_support_fun ()
```

7.26.3.25 likelihood() [1/4]

7.26.3.26 likelihood() [2/4]

7.26.3.27 likelihood() [3/4]

7.26.3.28 likelihood() [4/4]

7.26.3.29 likelihood_total()

7.26.3.30 nterms()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarrow
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
unsigned int Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >\leftarrow
::nterms ( ) const [noexcept]
```

7.26.3.31 operator=()

7.26.3.32 print()

```
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 >::print ()
const
```

Prints information about the model.

7.26.3.33 print_stats()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\longleftrightarrow Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::print_stats ( uint i ) const
```

7.26.3.34 sample() [1/2]

7.26.3.35 sample() [2/2]

7.26.3.36 set_counters()

7.26.3.37 set_keygen()

7.26.3.38 set_rengine()

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

7.26.3.39 set_rules()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::set_rules (
Rules< Array_Type, Data_Rule_Type > * rules_ )
```

7.26.3.40 set_rules_dyn()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_↔
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::set_rules_dyn
(
Rules< Array_Type, Data_Rule_Dyn_Type > * rules_ )
```

7.26.3.41 set_seed()

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

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

7.26.3.42 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	

7.26.3.43 size()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
unsigned int Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::size
( ) const [noexcept]
```

7.26.3.44 size_unique()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
unsigned int Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >\times
::size_unique ( ) const [noexcept]
```

7.26.3.45 store psets()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::store_psets (
) [noexcept]
```

7.26.3.46 support_size()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
unsigned int Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >\times
::support_size ( ) const [noexcept]
```

7.26.3.47 transform_model()

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

• include/barry/model-bones.hpp

7.27 NetCounterData Class Reference

Data class used to store arbitrary uint or double vectors.

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

Public Member Functions

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

Public Attributes

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

7.27.1 Detailed Description

Data class used to store arbitrary uint or double vectors.

Definition at line 56 of file network.hpp.

7.27.2 Constructor & Destructor Documentation

7.27.2.1 NetCounterData() [1/2]

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

Definition at line 62 of file network.hpp.

7.27.2.2 NetCounterData() [2/2]

Definition at line 63 of file network.hpp.

7.27.2.3 ∼NetCounterData()

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

Definition at line 68 of file network.hpp.

7.27.3 Member Data Documentation

7.27.3.1 indices

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

Definition at line 59 of file network.hpp.

7.27.3.2 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

7.28 NetworkData Class Reference

Data class for Networks.

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

Public Member Functions

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

Constructor using a single attribute.

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

Constructor using multiple attributes.

∼NetworkData ()

Public Attributes

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

7.28.1 Detailed Description

Data class for Networks.

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

7.28.2 Constructor & Destructor Documentation

7.28.2.1 NetworkData() [1/3]

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

Definition at line 25 of file network.hpp.

7.28.2.2 NetworkData() [2/3]

Constructor using a single attribute.

Parameters

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

Definition at line 33 of file network.hpp.

7.28.2.3 NetworkData() [3/3]

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

Constructor using multiple attributes.

7.29 Node Class Reference 165

Parameters

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

Definition at line 45 of file network.hpp.

7.28.2.4 ~NetworkData()

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

Definition at line 51 of file network.hpp.

7.28.3 Member Data Documentation

7.28.3.1 directed

bool NetworkData::directed = true

Definition at line 22 of file network.hpp.

7.28.3.2 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

7.29 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
- · unsigned int noffspring () const noexcept
- bool is_leaf () const noexcept

Construct a new Node object

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

Public Attributes

· unsigned int id

Id of the node (as specified in the input)

· unsigned int ord

Order in which the node was created.

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

Observed annotations (only defined for Geese)

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

Arrays given all possible states.

Node * parent = nullptr

Parent node.

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

Offspring nodes.

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

ID of the array in the model.

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

Induced subtree probabilities.

std::vector< double > probability

The probability of observing each state.

7.29.1 Detailed Description

A single node for the model.

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

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

7.29.2 Constructor & Destructor Documentation

7.29 Node Class Reference 167

7.29.2.1 Node() [1/5]

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

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

7.29.2.2 Node() [2/5]

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

Definition at line 56 of file geese-node-bones.hpp.

7.29.2.3 Node() [3/5]

Definition at line 62 of file geese-node-bones.hpp.

7.29.2.4 Node() [4/5]

```
Node::Node (
          Node && x ) [inline], [noexcept]
```

Definition at line 69 of file geese-node-bones.hpp.

7.29.2.5 Node() [5/5]

```
Node::Node ( {\tt const\ Node\ \&\ x\ )} \quad [{\tt inline}]
```

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

7.29.2.6 ∼Node()

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

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

7.29.3 Member Function Documentation

7.29.3.1 get_parent()

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

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

7.29.3.2 is_leaf()

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

Definition at line 109 of file geese-node-bones.hpp.

7.29.3.3 noffspring()

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

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

7.29.4 Member Data Documentation

7.29.4.1 annotations

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

Observed annotations (only defined for Geese)

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

7.29 Node Class Reference 169

7.29.4.2 array

phylocounters::PhyloArray Node::array

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

7.29.4.3 arrays

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

Arrays given all possible states.

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

7.29.4.4 duplication

bool Node::duplication

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

7.29.4.5 id

unsigned int Node::id

Id of the node (as specified in the input)

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

7.29.4.6 narray

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

ID of the array in the model.

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

7.29.4.7 offspring

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

Offspring nodes.

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

7.29.4.8 ord

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

Order in which the node was created.

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

7.29.4.9 parent

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

Parent node.

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

7.29.4.10 probability

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

The probability of observing each state.

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

7.29.4.11 subtree_prob

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

Induced subtree probabilities.

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

7.29.4.12 visited

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

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

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

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

7.30 NodeData Class Reference

Data definition for the PhyloArray class.

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

Public Member Functions

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

Public Attributes

```
std::vector< double > blengths = {}std::vector< bool > states = {}
```

• bool duplication = true

7.30.1 Detailed Description

Data definition for the PhyloArray class.

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

This holds basic information about a given node.

Definition at line 38 of file phylo.hpp.

7.30.2 Constructor & Destructor Documentation

7.30.2.1 NodeData()

Definition at line 58 of file phylo.hpp.

7.30.3 Member Data Documentation

7.30.3.1 blengths

```
std::vector< double > NodeData::blengths = {}
```

Branch length.

Definition at line 44 of file phylo.hpp.

7.30.3.2 duplication

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

Definition at line 54 of file phylo.hpp.

7.30.3.3 states

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

State of the parent node.

Definition at line 49 of file phylo.hpp.

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

• include/barry/counters/phylo.hpp

7.31 PhyloCounterData Class Reference

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

Public Member Functions

- PhyloCounterData (std::vector< uint > data_, std::vector< double > *counters_=nullptr)
- PhyloCounterData ()
- uint at (uint d)
- uint operator() (uint d)
- uint operator[] (uint d)
- void reserve (uint x)
- void push_back (uint x)
- void shrink_to_fit ()
- uint size ()
- std::vector< uint >::iterator begin ()
- std::vector< uint >::iterator end ()
- bool empty ()
- std::vector< double > * get_counters ()

7.31.1 Detailed Description

Definition at line 69 of file phylo.hpp.

7.31.2 Constructor & Destructor Documentation

7.31.2.1 PhyloCounterData() [1/2]

Definition at line 75 of file phylo.hpp.

7.31.2.2 PhyloCounterData() [2/2]

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

Definition at line 80 of file phylo.hpp.

7.31.3 Member Function Documentation

7.31.3.1 at()

Definition at line 82 of file phylo.hpp.

7.31.3.2 begin()

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

Definition at line 90 of file phylo.hpp.

7.31.3.3 empty()

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

Definition at line 93 of file phylo.hpp.

7.31.3.4 end()

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

Definition at line 91 of file phylo.hpp.

7.31.3.5 get_counters()

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

Definition at line 94 of file phylo.hpp.

7.31.3.6 operator()()

Definition at line 83 of file phylo.hpp.

7.31.3.7 operator[]()

```
uint PhyloCounterData::operator[] (
          uint d) [inline]
```

Definition at line 84 of file phylo.hpp.

7.31.3.8 push_back()

Definition at line 86 of file phylo.hpp.

7.31.3.9 reserve()

Definition at line 85 of file phylo.hpp.

7.31.3.10 shrink_to_fit()

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

Definition at line 87 of file phylo.hpp.

7.31.3.11 size()

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

Definition at line 88 of file phylo.hpp.

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

include/barry/counters/phylo.hpp

7.32 PhyloRuleDynData Class Reference

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

Public Member Functions

- PhyloRuleDynData (const std::vector< double > *counts_, uint pos_, uint lb_, uint ub_, uint duplication_)
- \sim PhyloRuleDynData ()

Public Attributes

- const std::vector< double > * counts
- uint pos
- uint lb
- uint ub
- · uint duplication

7.32.1 Detailed Description

Definition at line 2147 of file phylo.hpp.

7.32.2 Constructor & Destructor Documentation

7.32.2.1 PhyloRuleDynData()

Definition at line 2155 of file phylo.hpp.

7.32.2.2 ~PhyloRuleDynData()

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

Definition at line 2164 of file phylo.hpp.

7.32.3 Member Data Documentation

7.32.3.1 counts

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

Definition at line 2149 of file phylo.hpp.

7.32.3.2 duplication

```
uint PhyloRuleDynData::duplication
```

Definition at line 2153 of file phylo.hpp.

7.32.3.3 lb

```
uint PhyloRuleDynData::lb
```

Definition at line 2151 of file phylo.hpp.

7.32.3.4 pos

uint PhyloRuleDynData::pos

Definition at line 2150 of file phylo.hpp.

7.32.3.5 ub

uint PhyloRuleDynData::ub

Definition at line 2152 of file phylo.hpp.

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

• include/barry/counters/phylo.hpp

7.33 PowerSet< Array_Type, Data_Rule_Type > Class Template Reference

Powerset of a binary array.

#include <powerset-bones.hpp>

 $Collaboration\ diagram\ for\ PowerSet < Array_Type,\ Data_Rule_Type >:$



Public Member Functions

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

Construct and destroy a PowerSet object

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

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

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

- void add_rule (Rule < Array_Type, Data_Rule_Type > rule)
- void add_rule (Rule_fun_type< Array_Type, Data_Rule_Type > count_fun_, Data_Rule_Type data_)

Getter functions

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

Public Attributes

- Array_Type EmptyArray
- $std::vector < Array_Type > data$
- Rules
 Array_Type, Data_Rule_Type > * rules
- uint N
- uint M
- bool rules deleted = false
- std::vector < size_t > coordinates_free
- std::vector< size_t > coordinates_locked
- size_t n_free
- size_t n_locked

7.33.1 Detailed Description

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

Powerset of a binary array.

Template Parameters

Array_Type	
Data_Rule_Type	

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

7.33.2 Constructor & Destructor Documentation

7.33.2.1 PowerSet() [1/3]

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

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

7.33.2.2 PowerSet() [2/3]

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

7.33.2.3 PowerSet() [3/3]

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

7.33.2.4 ∼PowerSet()

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

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

7.33.3 Member Function Documentation

7.33.3.1 add_rule() [1/2]

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

7.33.3.2 add_rule() [2/2]

Definition at line 184 of file powerset-meat.hpp.

7.33.3.3 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 74 of file powerset-bones.hpp.

7.33.3.4 calc()

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

Definition at line 146 of file powerset-meat.hpp.

7.33.3.5 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 75 of file powerset-bones.hpp.

7.33.3.6 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 73 of file powerset-bones.hpp.

7.33.3.7 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 72 of file powerset-bones.hpp.

7.33.3.8 init support()

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

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

7.33.3.9 operator[]()

Definition at line 77 of file powerset-bones.hpp.

7.33.3.10 reset()

Definition at line 162 of file powerset-meat.hpp.

7.33.3.11 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 76 of file powerset-bones.hpp.

7.33.4 Member Data Documentation

7.33.4.1 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 32 of file powerset-bones.hpp.

7.33.4.2 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 33 of file powerset-bones.hpp.

7.33.4.3 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 25 of file powerset-bones.hpp.

7.33.4.4 EmptyArray

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

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

7.33.4.5 M

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

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

7.33.4.6 N

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

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

7.33.4.7 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 34 of file powerset-bones.hpp.

7.33.4.8 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 35 of file powerset-bones.hpp.

7.33.4.9 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 26 of file powerset-bones.hpp.

7.33.4.10 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 29 of file powerset-bones.hpp.

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

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

7.34 Progress Class Reference

A simple progress bar.

```
#include progress.hpp>
```

Public Member Functions

```
• Progress (int n_, int width_)
```

- ∼Progress ()
- void next ()
- void end ()

7.34.1 Detailed Description

A simple progress bar.

Definition at line 11 of file progress.hpp.

7.34.2 Constructor & Destructor Documentation

7.34.2.1 Progress()

```
Progress::Progress (
          int n_,
          int width_ ) [inline]
```

Definition at line 30 of file progress.hpp.

7.34.2.2 ∼Progress()

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

Definition at line 23 of file progress.hpp.

7.34.3 Member Function Documentation

7.34.3.1 end()

```
void Progress::end ( ) [inline]
```

Definition at line 52 of file progress.hpp.

7.34.3.2 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

7.35 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

- \sim Rule ()
- Data_Type & D ()

Read/Write access to the data.

bool operator() (const Array_Type &a, uint i, uint j)

Construct a new Rule object

Construct a new Rule object

Parameters

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

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

7.35.1 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 22 of file rules-bones.hpp.

7.35.2 Constructor & Destructor Documentation

7.35.2.1 Rule() [1/2]

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

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

7.35.2.2 Rule() [2/2]

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

7.35.2.3 ∼Rule()

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

Definition at line 47 of file rules-bones.hpp.

7.35.3 Member Function Documentation

7.35.3.1 D()

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

Read/Write access to the data.

7.35.3.2 operator()()

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

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

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

7.36 Rules < Array_Type, Data_Type > Class Template Reference

Vector of objects of class Rule.

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

Public Member Functions

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

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_)

7.36.1 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 62 of file rules-bones.hpp.

7.36.2 Constructor & Destructor Documentation

7.36.2.1 Rules() [1/2]

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

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

7.36.2.2 Rules() [2/2]

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

7.36.2.3 ∼Rules()

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

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

7.36.3 Member Function Documentation

7.36.3.1 add_rule() [1/2]

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

7.36.3.2 add_rule() [2/2]

Definition at line 57 of file rules-meat.hpp.

7.36.3.3 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 88 of file rules-meat.hpp.

7.36.3.4 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 72 of file rules-meat.hpp.

7.36.3.5 operator=()

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

7.36.3.6 size()

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

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

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

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

7.37 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 (uint i, uint j)

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

- void count current (uint i, uint i)
- 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

7.37.1 Detailed Description

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

Count stats for a single Array.

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

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

7.37.2 Constructor & Destructor Documentation

7.37.2.1 StatsCounter() [1/3]

Creator of a StatsCounter

Parameters

Array←	A const pointer to a BArray.

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

7.37.2.2 StatsCounter() [2/3]

Copy constructor.

Parameters

counter

7.37.2.3 StatsCounter() [3/3]

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

Can be created without setting the array.

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

7.37.2.4 ~StatsCounter()

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

7.37.3 Member Function Documentation

7.37.3.1 add_counter()

7.37.3.2 count_all()

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

Definition at line 102 of file statscounter-meat.hpp.

7.37.3.3 count_current()

7.37.3.4 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.

7.37.3.5 get_counters()

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

7.37.3.6 get_descriptions()

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

7.37.3.7 get_names()

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

7.37.3.8 reset_array()

Changes the reference array for the counting.

Parameters

Array←	A pointer to an array of class Array_Type.
1_	

7.37.3.9 set_counters()

7.37.3.10 size()

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

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

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

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

7.38 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 (uint N_, uint M_)

Constructor specifying the dimensions of the array (empty).

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

Computes the entire support.

std::vector< double > get_counts () const

- std::vector< double > * get_current_stats () List current statistics.
- · void print () const
- const FreqTable & 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.	1
1_		

- 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 uint &i, const uint &j)

Public Attributes

uint N

- uint M
- bool delete counters = true
- bool delete_rules = true
- bool delete_rules_dyn = true
- uint 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

7.38.1 Detailed Description

```
template < typename Array_Type = BArray <>>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
class 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 35 of file support-bones.hpp.

7.38.2 Constructor & Destructor Documentation

7.38.2.1 Support() [1/3]

Constructor passing a reference Array.

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

7.38.2.2 Support() [2/3]

Constructor specifying the dimensions of the array (empty).

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

7.38.2.3 Support() [3/3]

```
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 >::Support ( )
[inline]
```

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

7.38.2.4 ∼Support()

```
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 >::~Support ()
[inline]
```

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

7.38.3 Member Function Documentation

7.38.3.1 add_counter()

7.38.3.2 add_rule() [1/2]

7.38.3.3 add_rule() [2/2]

7.38.3.4 add_rule_dyn() [1/2]

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarrow Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::add_rule_\leftarrow dyn ( Rule< Array_Type, Data_Rule_Dyn_Type > * f_- )
```

7.38.3.5 add_rule_dyn() [2/2]

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarrow Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::add_rule_\leftarrow dyn ( Rule< Array_Type, Data_Rule_Dyn_Type > f_- )
```

7.38.3.6 calc()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::calc (
    std::vector< Array_Type > * array_bank = nullptr,
    std::vector< double > * stats_bank = nullptr,
    unsigned int max_num_elements_ = 0u )
```

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.

7.38.3.7 eval_rules_dyn()

7.38.3.8 get_counters()

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

Vector of couter functions.

7.38.3.9 get_counts()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< double > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn\times
_Type >::qet_counts ( ) const
```

7.38.3.10 get_current_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 >* Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_←
Dyn_Type >::get_current_stats ()
```

List current statistics.

7.38.3.11 get_data()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
const FreqTable& Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type
>::get_data ( ) const
```

7.38.3.12 get_rules()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Rules<Array_Type, Data_Rule_Type>* Support< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_rules ()
```

Vector of static rules (cells to iterate).

7.38.3.13 get_rules_dyn()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Rules<Array_Type, Data_Rule_Dyn_Type>* Support< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_rules_dyn ()
```

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

7.38.3.14 init support()

7.38.3.15 print()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::print ()
const
```

7.38.3.16 reset_array() [1/2]

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::reset_array
( )
```

7.38.3.17 reset_array() [2/2]

7.38.3.18 set counters()

7.38.3.19 set_rules()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::set_rules (
    Rules< Array_Type, Data_Rule_Type > * rules_ )
```

7.38.3.20 set_rules_dyn()

7.38.4 Member Data Documentation

7.38.4.1 change_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 > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn←
_Type >::change_stats
```

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

7.38.4.2 coordiantes_n_free

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
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 71 of file support-bones.hpp.

7.38.4.3 coordiantes n locked

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
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\times
n locked
```

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

7.38.4.4 coordinates_free

```
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 > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn←
_Type >::coordinates_free
```

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

7.38.4.5 coordinates_locked

```
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 > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn←
_Type >::coordinates_locked
```

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

7.38.4.6 current 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 > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn←
_Type >::current_stats
```

Definition at line 68 of file support-bones.hpp.

7.38.4.7 delete_counters

```
template<typename Array_Type = BArray<>, 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 62 of file support-bones.hpp.

7.38.4.8 delete rules

```
template<typename Array_Type = BArray<>, 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 63 of file support-bones.hpp.

7.38.4.9 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 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_←
rules_dyn = true
```

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

7.38.4.10 hashes

```
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 > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn←
_Type >::hashes
```

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

7.38.4.11 hashes initialized

```
template<typename Array_Type = BArray<>, 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 75 of file support-bones.hpp.

7.38.4.12 M

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

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

7.38.4.13 max_num_elements

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
uint Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::max_num_←
elements = BARRY_MAX_NUM_ELEMENTS
```

Definition at line 65 of file support-bones.hpp.

7.38.4.14 N

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

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

7.38.4.15 n counters

```
template<typename Array_Type = BArray<>, 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 76 of file support-bones.hpp.

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

• include/barry/support-bones.hpp

7.39 vecHasher < T > Struct Template Reference

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

Public Member Functions

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

7.39.1 Detailed Description

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

Definition at line 106 of file typedefs.hpp.

7.39.2 Member Function Documentation

7.39.2.1 operator()()

Definition at line 109 of file typedefs.hpp.

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

• include/barry/typedefs.hpp

Chapter 8

File Documentation

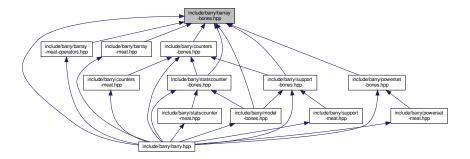
8.1 include/barry/barray-bones.hpp File Reference

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



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



Classes

class BArray < Cell_Type, Data_Type >
 Baseline class for binary arrays.

Macros

• #define BARRAY_BONES_HPP 1

8.1.1 Macro Definition Documentation

8.1.1.1 BARRAY_BONES_HPP

#define BARRAY_BONES_HPP 1

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

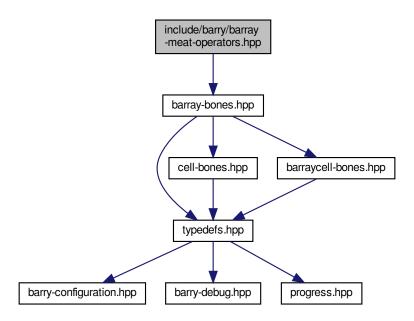
8.2 include/barry/barray-iterator.hpp File Reference

Classes

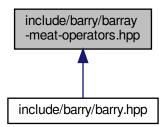
class ConstBArrayRowIter< Cell_Type, Data_Type >

8.3 include/barry/barray-meat-operators.hpp File Reference

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



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



Macros

- #define BARRY_BARRAY_MEAT_OPERATORS_HPP 1
- #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]

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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 (uint i=0u;i< nrow();++i) for(uint j=0u = el[POS(i, j)]
- j< ncol();++j) this-> operator() (i, j)+
- BARRAY_TEMPLATE (BARRAY_TYPE()&, operator+=)(const Cell_Type &rhs)
- BARRAY_TEMPLATE (BARRAY_TYPE()&, operator-=)(const BArray< Cell_Type
- BARRAY_TEMPLATE (BARRAY_TYPE()&, operator-=)(const Cell_Type &rhs)
- BARRAY TEMPLATE (BARRAY TYPE()&, operator*=)(const Cell Type &rhs)
- BARRAY_TEMPLATE (BARRAY_TYPE()&, operator/=)(const Cell_Type &rhs)

Variables

- Data_Type & rhs
- return * this

8.3.1 Macro Definition Documentation

8.3.1.1 BARRAY_TEMPLATE

Definition at line 11 of file barray-meat-operators.hpp.

8.3.1.2 BARRAY_TEMPLATE_ARGS

```
template BARRAY_TEMPLATE_ARGS() <typename Cell_Type, typename Data_Type>
```

Definition at line 9 of file barray-meat-operators.hpp.

8.3.1.3 BARRAY_TYPE

```
template Data_Type BARRAY_TYPE( ) BArray<Cell_Type, Data_Type>
```

Definition at line 7 of file barray-meat-operators.hpp.

8.3.1.4 BARRY_BARRAY_MEAT_OPERATORS_HPP

```
#define BARRY_BARRAY_MEAT_OPERATORS_HPP 1
```

Definition at line 5 of file barray-meat-operators.hpp.

8.3.1.5 COL

Definition at line 15 of file barray-meat-operators.hpp.

8.3.1.6 ROW

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

8.3.2 Function Documentation

8.3.2.1 BARRAY_TEMPLATE() [1/6]

Definition at line 88 of file barray-meat-operators.hpp.

8.3.2.2 BARRAY_TEMPLATE() [2/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator+ ) const
```

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8.3.2.3 BARRAY_TEMPLATE() [3/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator+ ) const &
```

Definition at line 46 of file barray-meat-operators.hpp.

8.3.2.4 BARRAY_TEMPLATE() [4/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator- ) const
```

8.3.2.5 BARRAY_TEMPLATE() [5/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator- ) const &
```

Definition at line 75 of file barray-meat-operators.hpp.

8.3.2.6 BARRAY_TEMPLATE() [6/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator/ ) const &
```

Definition at line 105 of file barray-meat-operators.hpp.

8.3.2.7 BARRAY_TEMPLATE_ARGS()

```
template BARRAY_TEMPLATE_ARGS ( ) const &
```

8.3.2.8 BARRAY_TYPE()

```
template const BARRAY_TYPE ( ) &
```

Definition at line 20 of file barray-meat-operators.hpp.

8.3.2.9 for()

```
for ( ) = el[POS(i, j)] [pure virtual]
```

Definition at line 66 of file barray-meat-operators.hpp.

8.3.2.10 operator()()

8.3.3 Variable Documentation

8.3.3.1 rhs

```
Data_Type & rhs
Initial value:
{
    checkdim_(*this, rhs)
```

Definition at line 33 of file barray-meat-operators.hpp.

8.3.3.2 this

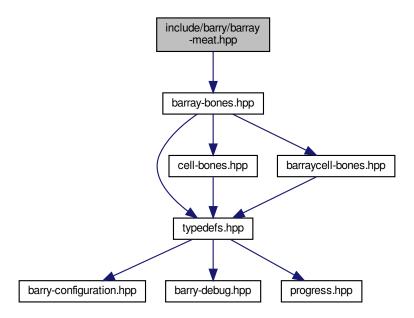
```
return * this
```

Definition at line 43 of file barray-meat-operators.hpp.

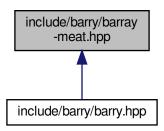
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8.4 include/barry/barray-meat.hpp File Reference

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



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



Macros

- #define BARRAY_TYPE() BArray<Cell_Type, Data_Type>
- #define BARRAY_TEMPLATE_ARGS() < typename Cell_Type, typename Data_Type>
- #define BARRAY_TEMPLATE(a, b) template BARRAY_TEMPLATE_ARGS() inline a BARRAY_TYPE()::b
- #define ROW(a) this->el_ij[a]
- #define COL(a) this->el_ji[a]

Functions

```
• BARRAY TEMPLATE (, BArray)(uint N_

    el_ij resize (N)

• el_ji resize (M)
• for (uint i=0u;i< source.size();++i)

    Data Type bool M (Array .M)

    BARRAY_TEMPLATE (BARRAY_TYPE() &, operator=)(const BArray< Cell_Type</li>

    BARRAY_TEMPLATE (, BArray)(BARRAY_TYPE() &&x) noexcept

    BARRAY TEMPLATE (BARRAY TYPE() &, operator=)(BARRAY TYPE() &&x) noexcept

• BARRAY TEMPLATE (bool, operator==)(const BARRAY TYPE() & Array )

    BARRAY_TEMPLATE (,~BArray)()

    BARRAY_TEMPLATE (void, set_data)(Data_Type *data_

    BARRAY TEMPLATE (Data Type *, D)()

• BARRAY_TEMPLATE (void, out_of_range)(uint i

    BARRAY_TEMPLATE (Cell_Type, get_cell)(uint i

    if (ROW(i).size()==0u) return(Cell Type) 0.0

• if (search !=ROW(i).end()) return search -> second.value
• return (Cell_Type) 0.0

    BARRAY_TEMPLATE (std::vector< Cell_Type >, get_row_vec)(uint i

    std::vector< Cell Type > ans (ncol(),(Cell Type) false)

    for (const auto &iter :row(i, false)) ans[iter.first]

    BARRAY_TEMPLATE (void, get_row_vec)(std

    BARRAY_TEMPLATE (BARRAY_TYPE() &, operator-=)(const std

    BARRAY_TEMPLATE (void, insert_cell)(uint i

• if (check_exists)

    COL (j).emplace(i

• & ROW (i)[j])

    BARRAY_TEMPLATE (void, swap_cells)(uint i0

if (report !=nullptr)(*report)
• if (check0 &check1)
• else if (!check0 &check1)

    else if (check0 &!check1)

    BARRAY_TEMPLATE (void, toggle_cell)(uint i

    BARRAY TEMPLATE (void, swap rows)(uint i0

• if (ROW(i0).size()==0u) move0
if (ROW(i1).size()==0u) move1
• if (!move0 &&!move1) return

    ROW (i0).swap(ROW(i1))

    BARRAY TEMPLATE (void, swap cols)(uint j0

if (COL(j0).size()==0u) check0
if (COL(j1).size()==0u) check1

    if (check0 &&check1)

• else if (check0 &&!check1)

    else if (!check0 &&check1)

    BARRAY_TEMPLATE (void, zero_row)(uint i

for (auto row=row0.begin();row !=row0.end();++row) rm_cell(i

    BARRAY TEMPLATE (void, zero col)(uint i

• if (COL(j).size()==0u) return

    BARRAY TEMPLATE (void, transpose)()

    BARRAY_TEMPLATE (void, clear)(bool hard)

    BARRAY_TEMPLATE (void, resize)(uint N_

• if (M_< M) for(uint j = N_
```

Variables

```
    uint M
```

- uint const std::vector< uint > & source
- uint const std::vector< uint > const std::vector< uint > & target
- uint const std::vector< uint > const std::vector< cell_Type > & value
- uint const std::vector< uint > const std::vector< Cell Type > bool add
- if(source.size() !=value.size()) throw std N = N_
- M = M
- return
- Data_Type & Array_
- Data_Type bool copy_data
- bool delete_data_
- data = data
- delete_data = delete_data_
- uint j const
- uint j
- auto search = ROW(i).find(j)
- · return ans
- uint const Cell
 Cell_Type > & v
- uint const Cell< Cell_Type > bool check_bounds
- uint const Cell
 Cell_Type > bool bool check_exists
- else
- NCells
- uint j0
- uint uint i1
- uint uint uint j1
- uint uint bool int int * report
- auto row0 = ROW(i)
- row first
- · row false
- auto col0 = COL(j)

8.4.1 Macro Definition Documentation

8.4.1.1 BARRAY_TEMPLATE

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

8.4.1.2 BARRAY_TEMPLATE_ARGS

```
#define BARRAY_TEMPLATE_ARGS( ) <typename Cell_Type, typename Data_Type>
```

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

8.4.1.3 BARRAY_TYPE

```
#define BARRAY_TYPE( ) BArray<Cell_Type, Data_Type>
```

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

8.4.1.4 COL

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

8.4.1.5 ROW

Definition at line 20 of file barray-meat.hpp.

8.4.2 Function Documentation

8.4.2.1 ans()

8.4.2.2 BARRAY_TEMPLATE() [1/23]

```
BARRAY_TEMPLATE (

BArray ) && [noexcept]
```

Definition at line 230 of file barray-meat.hpp.

8.4.2.3 BARRAY_TEMPLATE() [2/23]

```
BARRAY_TEMPLATE (
BArray )
```

8.4.2.4 BARRAY_TEMPLATE() [3/23]

```
BARRAY_TEMPLATE ( \sim \textit{BArray} \ )
```

Definition at line 339 of file barray-meat.hpp.

8.4.2.5 BARRAY_TEMPLATE() [4/23]

Definition at line 586 of file barray-meat.hpp.

8.4.2.6 BARRAY_TEMPLATE() [5/23]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE() & ,
          operator ) && [noexcept]
```

Definition at line 272 of file barray-meat.hpp.

8.4.2.7 BARRAY_TEMPLATE() [6/23]

8.4.2.8 BARRAY_TEMPLATE() [7/23]

```
BARRAY_TEMPLATE (
          bool ,
          operator = = ) const &
```

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

8.4.2.9 BARRAY_TEMPLATE() [8/23]

8.4.2.10 BARRAY_TEMPLATE() [9/23]

Definition at line 361 of file barray-meat.hpp.

8.4.2.11 BARRAY_TEMPLATE() [10/23]

```
BARRAY_TEMPLATE (
          std::vector< Cell_Type > ,
          get_row_vec )
```

8.4.2.12 BARRAY_TEMPLATE() [11/23]

```
BARRAY_TEMPLATE (
     void ,
     clear )
```

Definition at line 1119 of file barray-meat.hpp.

8.4.2.13 BARRAY_TEMPLATE() [12/23]

Definition at line 441 of file barray-meat.hpp.

8.4.2.14 BARRAY_TEMPLATE() [13/23]

8.4.2.15 BARRAY_TEMPLATE() [14/23]

8.4.2.16 BARRAY_TEMPLATE() [15/23]

```
BARRAY_TEMPLATE (
            void ,
            resize )
```

8.4.2.17 BARRAY_TEMPLATE() [16/23]

8.4.2.18 BARRAY_TEMPLATE() [17/23]

8.4.2.19 BARRAY_TEMPLATE() [18/23]

```
BARRAY_TEMPLATE (
     void ,
     swap_cols )
```

8.4.2.20 BARRAY_TEMPLATE() [19/23]

```
BARRAY_TEMPLATE (
     void ,
     swap_rows )
```

8.4.2.21 BARRAY_TEMPLATE() [20/23]

8.4.2.22 BARRAY_TEMPLATE() [21/23]

Definition at line 1058 of file barray-meat.hpp.

8.4.2.23 BARRAY_TEMPLATE() [22/23]

8.4.2.24 BARRAY_TEMPLATE() [23/23]

```
BARRAY_TEMPLATE (
     void ,
     zero_row )
```

8.4.2.25 COL()

```
COL (
```

```
8.4.2.26 for() [1/3]
```

```
for (
     auto row = row0.begin();row !=row0.end();++row )
```

8.4.2.27 for() [2/3]

8.4.2.28 for() [3/3]

```
for ( )
```

Definition at line 51 of file barray-meat.hpp.

8.4.2.29 if() [1/17]

```
else if (
    !check0 && check1 )
```

Definition at line 997 of file barray-meat.hpp.

8.4.2.30 if() [2/17]

```
else if (
    !check0 & check1 )
```

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

8.4.2.31 if() [3/17]

```
if (
    !move0 &&! move1 )
```

8.4.2.32 if() [4/17]

Definition at line 853 of file barray-meat.hpp.

```
8.4.2.33 if() [5/17]
```

Definition at line 988 of file barray-meat.hpp.

8.4.2.34 if() [6/17]

```
if ( check0 && check1)
```

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

8.4.2.35 if() [7/17]

```
if ( check0 & check1)
```

Definition at line 827 of file barray-meat.hpp.

8.4.2.36 if() [8/17]

```
else if (
          check_exists = = CHECK::BOTH )
```

Definition at line 668 of file barray-meat.hpp.

8.4.2.37 if() [9/17]

```
if ( \label{eq:col} \mathtt{COL}(\mathtt{j}).\mathtt{size}(\mathtt{)} \ = = 0u \ \mathtt{)}
```

```
8.4.2.38 if() [10/17]
```

```
if ( COL(j0).size() = =0u )
```

8.4.2.39 if() [11/17]

```
if ( \label{eq:col} \mathtt{COL}(\mathtt{j1}).\mathtt{size}() \ = = 0u \ )
```

8.4.2.40 if() [12/17]

```
else if ( ) = N_
```

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

8.4.2.41 if() [13/17]

```
if (
    report ! = nullptr )
```

8.4.2.42 if() [14/17]

```
if ( \label{eq:row_row_row} \mbox{ROW(i).size()} \ = \ = 0 \mbox{$u$} \ )
```

8.4.2.43 if() [15/17]

```
if ( \label{eq:row_row} \mbox{ROW(i0).size()} \ = \ = \mbox{$0$u } \mbox{)}
```

8.4.2.44 if() [16/17]

```
if ( \label{eq:row_row_row} \text{ROW(i1).size()} \quad = = 0u \text{ )}
```

```
8.4.2.45 if() [17/17]
```

```
if (
    search ! = ROW(i).end() ) -> second.value
```

8.4.2.46 M()

```
Data_Type bool M ( \label{eq:continuous} {\tt Array\_.} \quad {\tt M} \; )
```

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

8.4.2.47 resize() [1/2]

```
el_ji resize (
M )
```

8.4.2.48 resize() [2/2]

```
el_ij resize (
N )
```

8.4.2.49 return()

8.4.2.50 ROW() [1/2]

```
& ROW ( i )
```

8.4.2.51 ROW() [2/2]

```
ROW ( i0 )
```

8.4.3 Variable Documentation

8.4.3.1 add

```
uint const std::vector< uint > const std::vector< uint > bool add

Initial value:
{
    if (source.size() != target.size())
        throw std::length_error("-source- and -target- don't match on length.")
```

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

8.4.3.2 ans

```
return ans
```

Definition at line 438 of file barray-meat.hpp.

8.4.3.3 Array_

```
Data_Type & Array_
```

Definition at line 134 of file barray-meat.hpp.

8.4.3.4 check_bounds

```
Initial value:
{
    if (check_bounds) {
       out_of_range(i0,0u);
       out_of_range(i1,0u);
    }
    bool move0=true, move1=true
```

Definition at line 661 of file barray-meat.hpp.

8.4.3.5 check_exists

```
uint bool int check_exists
Initial value:
{
    if (check_bounds)
        out_of_range(i,j)
```

Definition at line 662 of file barray-meat.hpp.

8.4.3.6 col0

```
auto col0 = COL(j)
```

Definition at line 1050 of file barray-meat.hpp.

8.4.3.7 const

```
uint bool check_bounds const

Initial value:
{
    if (i >= N)
        throw std::range_error("The row is out of range.")
```

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

8.4.3.8 copy_data

```
Data_Type bool copy_data
```

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

8.4.3.9 data

```
data = data_
```

Definition at line 354 of file barray-meat.hpp.

8.4.3.10 delete_data

```
delete_data = delete_data_
```

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

8.4.3.11 delete_data_

Definition at line 348 of file barray-meat.hpp.

8.4.3.12 else

Definition at line 692 of file barray-meat.hpp.

8.4.3.13 false

row false

Definition at line 1031 of file barray-meat.hpp.

8.4.3.14 first

row first

Definition at line 1031 of file barray-meat.hpp.

8.4.3.15 i1

```
uint i1
```

Definition at line 765 of file barray-meat.hpp.

8.4.3.16 j

```
uint j
```

Initial value:

```
if (init_fun == nullptr)
    return 0.0
```

Definition at line 403 of file barray-meat.hpp.

8.4.3.17 j0

```
uint j0
```

Definition at line 764 of file barray-meat.hpp.

8.4.3.18 j1

```
uint j1
```

Definition at line 765 of file barray-meat.hpp.

8.4.3.19 M

```
M = M_{\underline{}}
```

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

8.4.3.20 M_

```
uint M_
Initial value:
{

    if (N_ < N)
        for (uint i = N_; i < N; ++i)
            zero_row(i, false)</pre>
```

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

8.4.3.21 N

```
if (source.size() != target.size()) throw std if (source.size() != value.size()) throw std N=N
```

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

8.4.3.22 NCells

NCells

Definition at line 696 of file barray-meat.hpp.

8.4.3.23 report

```
uint uint uint bool int int* report
```

Definition at line 768 of file barray-meat.hpp.

8.4.3.24 return

return

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

8.4.3.25 row0

```
auto row0 = ROW(i)
```

Definition at line 1029 of file barray-meat.hpp.

8.4.3.26 search

```
auto search = ROW(i).find(j)
```

Definition at line 415 of file barray-meat.hpp.

8.4.3.27 source

```
uint const std::vector< uint > & source
```

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

8.4.3.28 target

```
uint const std::vector< uint > const std::vector< uint > & target
```

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

8.4.3.29 v

```
uint Cell_Type v
```

Definition at line 660 of file barray-meat.hpp.

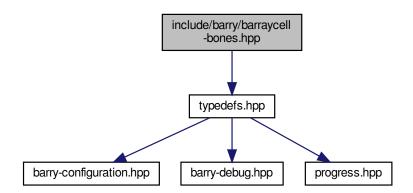
8.4.3.30 value

uint const std::vector< uint > const std::vector< uint > const std::vector< Cell_Type > & value

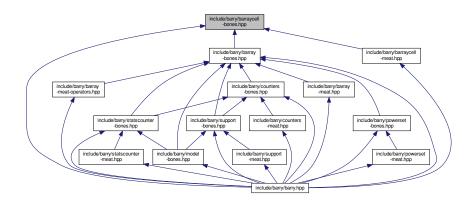
Definition at line 33 of file barray-meat.hpp.

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

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



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



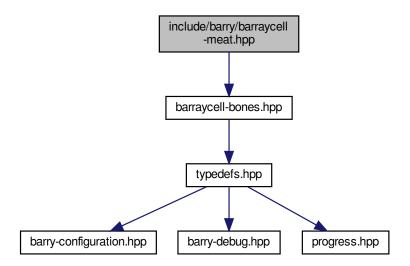
Classes

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

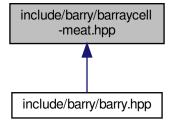
8.6 include/barry/barraycell-meat.hpp File Reference

#include "barraycell-bones.hpp"

Include dependency graph for barraycell-meat.hpp:



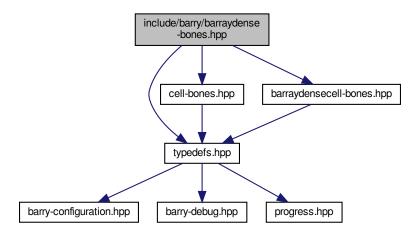
This graph shows which files directly or indirectly include this file:



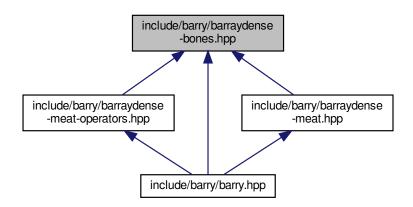
8.7 include/barry/barraydense-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "cell-bones.hpp"
#include "barraydensecell-bones.hpp"
```

Include dependency graph for barraydense-bones.hpp:



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



Classes

class BArrayDense < Cell_Type, Data_Type >
 Baseline class for binary arrays.

Macros

• #define BARRY_BARRAYDENSE_BONES_HPP 1

8.7.1 Macro Definition Documentation

8.7.1.1 BARRY_BARRAYDENSE_BONES_HPP

#define BARRY_BARRAYDENSE_BONES_HPP 1

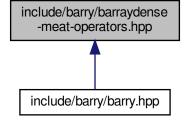
Definition at line 8 of file barraydense-bones.hpp.

8.8 include/barry/barraydense-meat-operators.hpp File Reference

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



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



Macros

- #define BARRY_BARRAYDENSE_MEAT_OPERATORS_HPP 1
- #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)

8.8.1 Macro Definition Documentation

8.8.1.1 BARRY_BARRAYDENSE_MEAT_OPERATORS_HPP

```
#define BARRY_BARRAYDENSE_MEAT_OPERATORS_HPP 1
```

Definition at line 5 of file barraydense-meat-operators.hpp.

8.8.1.2 BDENSE_TEMPLATE

Definition at line 11 of file barraydense-meat-operators.hpp.

8.8.1.3 BDENSE_TEMPLATE_ARGS

```
template BDENSE_TEMPLATE_ARGS() <typename Cell_Type, typename Data_Type>
```

Definition at line 9 of file barraydense-meat-operators.hpp.

8.8.1.4 BDENSE_TYPE

```
template Data_Type BDENSE_TYPE() BArrayDense<Cell_Type, Data_Type>
```

Definition at line 7 of file barraydense-meat-operators.hpp.

8.8.1.5 COL

Definition at line 15 of file barraydense-meat-operators.hpp.

8.8.1.6 POS

```
#define POS( \label{eq:a_b} a, \\ b \ ) \ (b)*N + (a)
```

Definition at line 16 of file barraydense-meat-operators.hpp.

8.8.1.7 POS N

Definition at line 17 of file barraydense-meat-operators.hpp.

8.8.1.8 ROW

Definition at line 14 of file barraydense-meat-operators.hpp.

8.8.2 Function Documentation

8.8.2.1 BDENSE_TEMPLATE() [1/4]

Definition at line 90 of file barraydense-meat-operators.hpp.

8.8.2.2 BDENSE_TEMPLATE() [2/4]

Definition at line 34 of file barraydense-meat-operators.hpp.

8.8.2.3 BDENSE_TEMPLATE() [3/4]

Definition at line 61 of file barraydense-meat-operators.hpp.

8.8.2.4 BDENSE_TEMPLATE() [4/4]

Definition at line 101 of file barraydense-meat-operators.hpp.

8.8.2.5 BDENSE_TEMPLATE_ARGS()

```
template BDENSE_TEMPLATE_ARGS ( ) const &
```

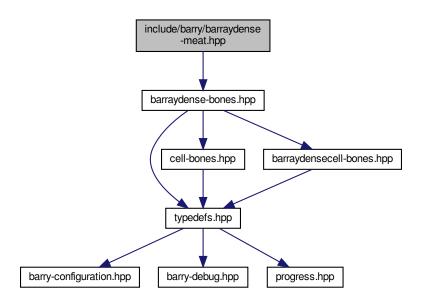
8.8.2.6 BDENSE_TYPE()

```
template const BDENSE_TYPE ( ) &
```

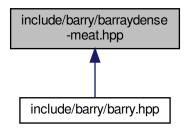
Definition at line 22 of file barraydense-meat-operators.hpp.

8.9 include/barry/barraydense-meat.hpp File Reference

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



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



Macros

- #define BDENSE_TYPE() BArrayDense<Cell_Type, Data_Type>
- #define BDENSE_TEMPLATE_ARGS() < typename Cell_Type, typename Data_Type>
- #define BDENSE_TEMPLATE(a, b) template BDENSE_TEMPLATE_ARGS() inline a BDENSE_TYPE()::b
- #define ROW(a) this->el_ij[a]
- #define COL(a) this->el_ji[a]
- #define POS(a, b) (b)*N + (a)
- #define POS_N(a, b, c) (b)*(c) + (a)
- #define ZERO_CELL static_cast<Cell_Type>(0.0)

Functions

```
    BDENSE_TEMPLATE (, BArrayDense)(uint N_

    el resize (N *M, ZERO_CELL)

• el_rowsums resize (N, ZERO_CELL)
• el colsums resize (M, ZERO CELL)

    for (uint i=0u;i< source.size();++i)</li>

    BDENSE_TEMPLATE (, BArrayDense)(const BDENSE_TYPE() &Array_

    bool M (Array .M)

• BDENSE_TEMPLATE (BDENSE_TYPE() &, operator=)(const BDENSE_TYPE() &Array_)
• BDENSE TEMPLATE (, BArrayDense)(BDENSE TYPE() &&x) noexcept

    BDENSE TEMPLATE (BDENSE TYPE() &, operator=)(BDENSE TYPE() &&x) noexcept

• BDENSE_TEMPLATE (bool, operator==)(const BDENSE_TYPE() &Array_)

    BDENSE TEMPLATE (, ~BArrayDense)()

    BDENSE_TEMPLATE (void, set_data)(Data_Type *data_

• BDENSE_TEMPLATE (Data_Type *, D)()
• BDENSE TEMPLATE (const Data Type *, D)() const
• BDENSE TEMPLATE (void, out of range)(uint i

    BDENSE TEMPLATE (Cell Type, get cell)(uint i

• BDENSE_TEMPLATE (std::vector< Cell_Type >, get_row_vec)(uint i

    std::vector< Cell Type > ans (ncol(), static cast< Cell Type >(false))

• BDENSE_TEMPLATE (void, get_row_vec)(std
• BDENSE TEMPLATE (Entries < Cell Type >, get entries)() const
• BDENSE TEMPLATE (bool, is empty)(uint i

    BDENSE TEMPLATE (unsigned int, nrow)() const noexcept

    BDENSE_TEMPLATE (unsigned int, ncol)() const noexcept

    BDENSE_TEMPLATE (unsigned int, nnozero)() const noexcept

• BDENSE TEMPLATE (Cell< Cell Type >, default val)() const
• BDENSE TEMPLATE (BDENSE TYPE() &, operator+=)(const std
• BDENSE TEMPLATE (BDENSE TYPE() &, operator-=)(const std

    BDENSE TEMPLATE (void, insert cell)(uint i

if (el[POS(i, j)]==BARRY_ZERO_DENSE)

    BDENSE_TEMPLATE (void, swap_cells)(uint i0

• if ((i0==i1) &&(j0==j1)) return
• rm cell (i0, j0, false, false)
• rm_cell (i1, j1, false, false)
• insert cell (i0, j0, val1, false, false)

    insert cell (i1, j1, val0, false, false)

• BDENSE_TEMPLATE (void, toggle_cell)(uint i
• else rm cell (i, j, false, false)
• BDENSE TEMPLATE (void, swap rows)(uint i0

    BDENSE TEMPLATE (void, swap cols)(uint j0

• BDENSE_TEMPLATE (void, zero_row)(uint i
• if (el rowsums[i]==ZERO CELL) return
• BDENSE_TEMPLATE (void, zero_col)(uint j
• if (el colsums[i]==ZERO CELL) return

    BDENSE TEMPLATE (void, transpose)()

• BDENSE TEMPLATE (void, clear)(bool hard)

    BDENSE TEMPLATE (void, resize)(uint N

• el resize (N_ *M_, ZERO_CELL)
• el rowsums resize (N, ZERO CELL)
• el colsums resize (M, ZERO CELL)
• BDENSE TEMPLATE (void, reserve)()
```

• BDENSE_TEMPLATE (void, print)(const char *fmt

va_start (args, fmt)

```
vprintf (fmt, args)
va_end (args)
BDENSE_TEMPLATE (const std::vector< Cell_Type > &, get_data)() const
BDENSE_TEMPLATE (const Cell_Type, rowsum)(unsigned int i) const
BDENSE_TEMPLATE (const Cell_Type, colsum)(unsigned int j) const
```

Variables

```
    uint M

• uint const std::vector< uint > & source
• uint const std::vector< uint > const std::vector< uint > & target

    uint const std::vector< uint > const std::vector< uint > const std::vector< Cell_Type > & value

    uint const std::vector< uint > const std::vector< cell_Type > bool add

    if(source.size() !=value.size()) throw std N = N_

    M = M_

    return

· bool copy_data
· bool delete_data_
• data = data_
delete_data = delete_data_
· uint j const
uint j
• return el [POS(i, j)] == ZERO_CELL
· return ans

    uint const Cell
    Cell_Type > & v

    uint const Cell
    Cell Type > bool check bounds

    uint const Cell
    Cell_Type > bool bool check_exists

el_rowsums [i] = (v.value - old)
• el_colsums [j] = (v.value - old)
uint j0
· uint uint i1
• uint uint uint j1
• uint uint uint bool int int * report
• Cell Type val0 = el[POS(i0,j0)]
• Cell_Type val1 = el[POS(i1,j1)]

    false

col
```

8.9.1 Macro Definition Documentation

8.9.1.1 BDENSE_TEMPLATE

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

8.9.1.2 BDENSE_TEMPLATE_ARGS

```
#define BDENSE_TEMPLATE_ARGS( ) <typename Cell_Type, typename Data_Type>
```

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

8.9.1.3 BDENSE_TYPE

```
#define BDENSE_TYPE( ) BArrayDense<Cell_Type, Data_Type>
```

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

8.9.1.4 COL

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

8.9.1.5 POS

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

Definition at line 32 of file barraydense-meat.hpp.

8.9.1.6 POS_N

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

8.9.1.7 ROW

Definition at line 30 of file barraydense-meat.hpp.

8.9.1.8 ZERO_CELL

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

Definition at line 38 of file barraydense-meat.hpp.

8.9.2 Function Documentation

8.9.2.1 ans()

8.9.2.2 BDENSE_TEMPLATE() [1/37]

Definition at line 240 of file barraydense-meat.hpp.

8.9.2.3 BDENSE_TEMPLATE() [2/37]

8.9.2.4 BDENSE_TEMPLATE() [3/37]

```
BDENSE_TEMPLATE (
BArrayDense )
```

8.9.2.5 BDENSE_TEMPLATE() [4/37]

```
BDENSE_TEMPLATE ( \sim BArrayDense )
```

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

8.9.2.6 BDENSE_TEMPLATE() [5/37]

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

8.9.2.7 BDENSE_TEMPLATE() [6/37]

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

8.9.2.8 BDENSE_TEMPLATE() [7/37]

Definition at line 257 of file barraydense-meat.hpp.

8.9.2.9 BDENSE_TEMPLATE() [8/37]

Definition at line 194 of file barraydense-meat.hpp.

8.9.2.10 BDENSE_TEMPLATE() [9/37]

```
BDENSE_TEMPLATE (
          bool ,
          is_empty )
```

8.9.2.11 BDENSE_TEMPLATE() [10/37]

```
BDENSE_TEMPLATE (
          bool ,
          operator = = ) const &
```

Definition at line 300 of file barraydense-meat.hpp.

8.9.2.12 BDENSE_TEMPLATE() [11/37]

Definition at line 554 of file barraydense-meat.hpp.

8.9.2.13 BDENSE_TEMPLATE() [12/37]

8.9.2.14 BDENSE_TEMPLATE() [13/37]

Definition at line 991 of file barraydense-meat.hpp.

8.9.2.15 BDENSE_TEMPLATE() [14/37]

Definition at line 986 of file barraydense-meat.hpp.

8.9.2.16 BDENSE_TEMPLATE() [15/37]

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

8.9.2.17 BDENSE_TEMPLATE() [16/37]

Definition at line 981 of file barraydense-meat.hpp.

8.9.2.18 BDENSE_TEMPLATE() [17/37]

Definition at line 341 of file barraydense-meat.hpp.

8.9.2.19 BDENSE_TEMPLATE() [18/37]

Definition at line 494 of file barraydense-meat.hpp.

8.9.2.20 BDENSE_TEMPLATE() [19/37]

```
BDENSE_TEMPLATE (
          std::vector< Cell_Type > ,
          get_row_vec )
```

8.9.2.21 BDENSE_TEMPLATE() [20/37]

```
BDENSE_TEMPLATE (
         unsigned int ,
         ncol ) const [noexcept]
```

Definition at line 540 of file barraydense-meat.hpp.

8.9.2.22 BDENSE_TEMPLATE() [21/37]

```
BDENSE_TEMPLATE (
          unsigned int ,
          nnozero ) const [noexcept]
```

Definition at line 544 of file barraydense-meat.hpp.

8.9.2.23 BDENSE_TEMPLATE() [22/37]

```
BDENSE_TEMPLATE (
         unsigned int ,
         nrow ) const [noexcept]
```

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

8.9.2.24 BDENSE_TEMPLATE() [23/37]

```
BDENSE_TEMPLATE (
     void ,
     clear )
```

Definition at line 888 of file barraydense-meat.hpp.

8.9.2.25 BDENSE_TEMPLATE() [24/37]

Definition at line 394 of file barraydense-meat.hpp.

8.9.2.26 BDENSE_TEMPLATE() [25/37]

8.9.2.27 BDENSE_TEMPLATE() [26/37]

8.9.2.28 BDENSE_TEMPLATE() [27/37]

8.9.2.29 BDENSE_TEMPLATE() [28/37]

```
BDENSE_TEMPLATE (
     void ,
     reserve )
```

Definition at line 938 of file barraydense-meat.hpp.

8.9.2.30 BDENSE_TEMPLATE() [29/37]

```
BDENSE_TEMPLATE (
     void ,
     resize )
```

8.9.2.31 BDENSE_TEMPLATE() [30/37]

8.9.2.32 BDENSE_TEMPLATE() [31/37]

8.9.2.33 BDENSE_TEMPLATE() [32/37]

8.9.2.34 BDENSE_TEMPLATE() [33/37]

```
BDENSE_TEMPLATE (
     void ,
     swap_rows )
```

8.9.2.35 BDENSE_TEMPLATE() [34/37]

8.9.2.36 BDENSE_TEMPLATE() [35/37]

```
BDENSE_TEMPLATE (
     void ,
     transpose )
```

Definition at line 860 of file barraydense-meat.hpp.

8.9.2.37 BDENSE_TEMPLATE() [36/37]

```
BDENSE_TEMPLATE (
            void ,
            zero_col )
```

8.9.2.38 BDENSE_TEMPLATE() [37/37]

```
BDENSE_TEMPLATE (
     void ,
     zero_row )
```

8.9.2.39 for()

```
for ( )
```

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

```
8.9.2.40 if() [1/4]
```

```
if (  ({\tt i0==i1)} \ \&\&\,({\tt j0==j1}) \ )
```

8.9.2.41 if() [2/4]

Definition at line 655 of file barraydense-meat.hpp.

```
8.9.2.42 if() [3/4]
```

```
if (
    el_colsums [j] = =ZERO_CELL )
```

8.9.2.43 if() [4/4]

8.9.2.44 insert_cell() [1/2]

8.9.2.45 insert_cell() [2/2]

8.9.2.46 M()

```
bool M ( \label{eq:Array_.} \text{Array}\_. \quad \textit{M} \ )
```

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

8.9.2.47 resize() [1/6]

8.9.2.48 resize() [2/6]

8.9.2.49 resize() [3/6]

```
el resize ( \label{eq:N*M, ZERO_CELL} \mbox{N * $M$,}
```

8.9.2.50 resize() [4/6]

```
el_rowsums resize (
          N ,
          ZERO_CELL )
```

8.9.2.51 resize() [5/6]

```
el resize ( \label{eq:n_* M_*, ZERO_CELL} N\_ * M\_,
```

8.9.2.52 resize() [6/6]

8.9.2.53 rm_cell() [1/3]

8.9.2.54 rm_cell() [2/3]

8.9.2.55 rm_cell() [3/3]

8.9.2.56 va_end()

```
va_end (
          args )
```

8.9.2.57 va_start()

8.9.2.58 vprintf()

8.9.3 Variable Documentation

8.9.3.1 add

```
uint const std::vector< uint > const std::vector< uint > bool add

Initial value:
{
    if (source.size() != target.size())
        throw std::length_error("-source- and -target- don't match on length.")
```

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

8.9.3.2 ans

```
return ans
```

Definition at line 390 of file barraydense-meat.hpp.

8.9.3.3 check_bounds

```
bool check_bounds

Initial value:
{
    if (check_bounds)
        out_of_range(i0,0u);
        out_of_range(i1,0u);
    }

for (uint j = 0u; j < M; ++j)
        std::swap(el[POS(i0, j)], el[POS(i1, j)])</pre>
```

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

8.9.3.4 check exists

```
uint bool int check_exists
Initial value:
{
    if (check_bounds)
        out_of_range(i,j)
```

Definition at line 647 of file barraydense-meat.hpp.

8.9.3.5 col

col

Definition at line 835 of file barraydense-meat.hpp.

8.9.3.6 const

const

Initial value:

```
if (i >= N)
    throw std::range_error("The row is out of range.")
```

Definition at line 352 of file barraydense-meat.hpp.

8.9.3.7 copy_data

bool copy_data

Definition at line 156 of file barraydense-meat.hpp.

8.9.3.8 data

```
data = data_
```

Definition at line 334 of file barraydense-meat.hpp.

8.9.3.9 delete_data

```
delete_data = delete_data_
```

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

8.9.3.10 delete_data_

Definition at line 328 of file barraydense-meat.hpp.

8.9.3.11 el

```
return el == ZERO_CELL
```

Definition at line 373 of file barraydense-meat.hpp.

8.9.3.12 el_colsums

```
el_colsums[j] = (v.value - old)
```

Definition at line 667 of file barraydense-meat.hpp.

8.9.3.13 el_rowsums

```
el_rowsums[i] = (v.value - old)
```

Definition at line 666 of file barraydense-meat.hpp.

8.9.3.14 else

Definition at line 662 of file barraydense-meat.hpp.

8.9.3.15 false

false

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

8.9.3.16 i1

uint i1

Definition at line 713 of file barraydense-meat.hpp.

8.9.3.17 j

j

Definition at line 365 of file barraydense-meat.hpp.

8.9.3.18 j0

uint j0

Definition at line 712 of file barraydense-meat.hpp.

8.9.3.19 j1

uint j1

Definition at line 713 of file barraydense-meat.hpp.

8.9.3.20 M

 $M = M_{\underline{}}$

Definition at line 57 of file barraydense-meat.hpp.

8.9.3.21 M_

```
uint M_
Initial value:
{
    std::vector< Cell_Type > el_tmp(el)
```

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

8.9.3.22 N

```
N = N_
```

Definition at line 56 of file barraydense-meat.hpp.

8.9.3.23 report

```
uint uint uint bool int int* report

Initial value:
{
    if (check_bounds) {
        out_of_range(i0, j0);
        out_of_range(i1, j1);
    }

    if (report != nullptr)
        (*report) = EXISTS::BOTH
```

Definition at line 716 of file barraydense-meat.hpp.

8.9.3.24 return

return

Definition at line 94 of file barraydense-meat.hpp.

8.9.3.25 source

```
uint const std::vector< uint >& source
```

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

8.9.3.26 target

```
uint const std::vector< uint > const std::vector< uint >& target
```

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

8.9.3.27 v

```
uint Cell_Type v
```

Definition at line 645 of file barraydense-meat.hpp.

8.9.3.28 val0

```
Cell_Type val0 = el[POS(i0,j0)]
```

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

8.9.3.29 val1

```
Cell_Type val1 = el[POS(i1, j1)]
```

Definition at line 735 of file barraydense-meat.hpp.

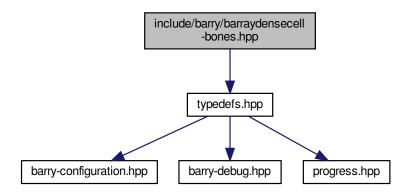
8.9.3.30 value

```
uint const std::vector< uint > const std::vector< cell_Type >&
value
```

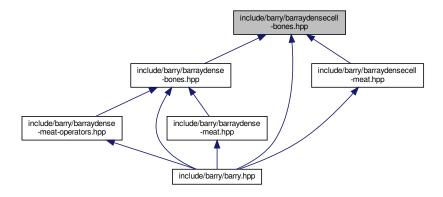
Definition at line 46 of file barraydense-meat.hpp.

8.10 include/barry/barraydensecell-bones.hpp File Reference

#include "typedefs.hpp"
Include dependency graph for barraydensecell-bones.hpp:



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

8.10.1 Macro Definition Documentation

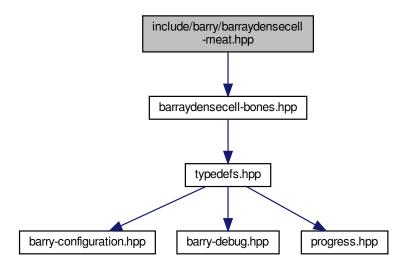
8.10.1.1 POS

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

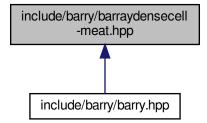
Definition at line 6 of file barraydensecell-bones.hpp.

8.11 include/barry/barraydensecell-meat.hpp File Reference

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



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



Macros

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

8.11.1 Macro Definition Documentation

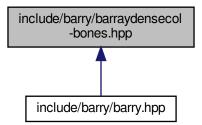
8.11.1.1 POS

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

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

8.12 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)

8.12.1 Macro Definition Documentation

8.12.1.1 POS

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

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

8.12.1.2 POS_N

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

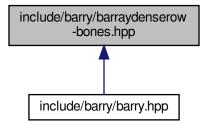
8.12.1.3 ZERO_CELL

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

Definition at line 6 of file barraydensecol-bones.hpp.

8.13 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)
```

8.13.1 Macro Definition Documentation

8.13.1.1 POS

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

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

8.13.1.2 POS N

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

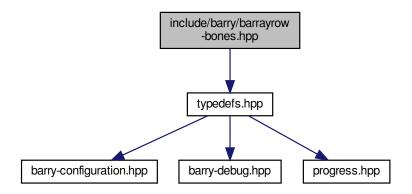
8.13.1.3 ZERO_CELL

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

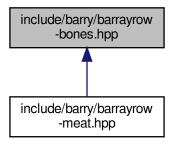
Definition at line 6 of file barraydenserow-bones.hpp.

8.14 include/barry/barrayrow-bones.hpp File Reference

#include "typedefs.hpp"
Include dependency graph for barrayrow-bones.hpp:



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



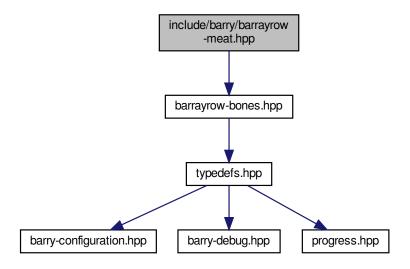
Classes

- class BArrayRow< Cell_Type, Data_Type >
- class BArrayRow_const< Cell_Type, Data_Type >

8.15 include/barry/barrayrow-meat.hpp File Reference

#include "barrayrow-bones.hpp"

Include dependency graph for barrayrow-meat.hpp:



Macros

- #define BARRY_BARRAYROW_MEAT_HPP 1
- #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)

8.15.1 Macro Definition Documentation

8.15.1.1 BARRY BARRAYROW MEAT HPP

#define BARRY_BARRAYROW_MEAT_HPP 1

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

8.15.1.2 BROW_TEMPLATE

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

8.15.1.3 BROW_TEMPLATE_ARGS

```
#define BROW_TEMPLATE_ARGS() <typename Cell_Type, typename Data_Type>
```

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

8.15.1.4 BROW_TYPE

```
#define BROW_TYPE( ) BArrayRow<Cell_Type, Data_Type>
```

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

8.15.2 Function Documentation

8.15.2.1 BROW_TEMPLATE() [1/5]

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

8.15.2.2 BROW_TEMPLATE() [2/5]

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

8.15.2.3 BROW_TEMPLATE() [3/5]

Definition at line 36 of file barrayrow-meat.hpp.

8.15.2.4 BROW_TEMPLATE() [4/5]

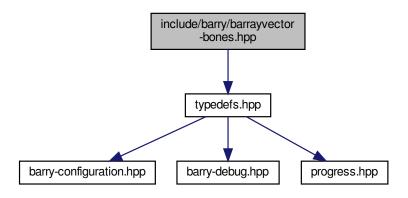
Definition at line 57 of file barrayrow-meat.hpp.

8.15.2.5 BROW_TEMPLATE() [5/5]

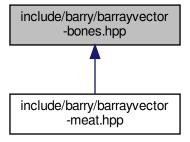
Definition at line 13 of file barrayrow-meat.hpp.

8.16 include/barry/barrayvector-bones.hpp File Reference

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



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

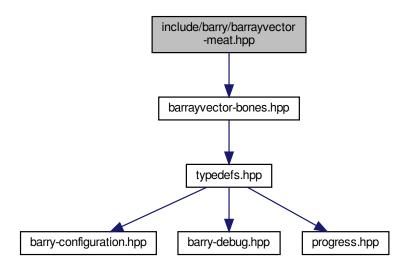


Classes

- class BArrayVector< Cell_Type, Data_Type >
 Row or column of a BArray
- class BArrayVector_const< Cell_Type, Data_Type >

8.17 include/barry/barrayvector-meat.hpp File Reference

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



Macros

• #define BARRY BARRAYVECTOR MEAT HPP 1

8.17.1 Macro Definition Documentation

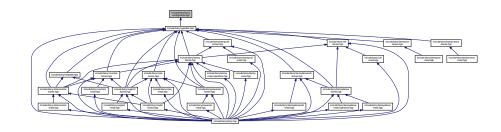
8.17.1.1 BARRY_BARRAYVECTOR_MEAT_HPP

```
#define BARRY_BARRAYVECTOR_MEAT_HPP 1
```

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

8.18 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 >(UINT_MAX/2u)
- template<typename Ta , typename Tb >
 using Map = std::map< Ta, Tb >

8.18.1 Macro Definition Documentation

8.18.1.1 BARRY_CHECK_SUPPORT

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

Definition at line 47 of file barry-configuration.hpp.

8.18.1.2 BARRY_ISFINITE

Definition at line 40 of file barry-configuration.hpp.

8.18.1.3 BARRY_MAX_NUM_ELEMENTS

```
#define BARRY_MAX_NUM_ELEMENTS static_cast< size_t >(UINT_MAX/2u)
```

Definition at line 55 of file barry-configuration.hpp.

8.18.1.4 BARRY_SAFE_EXP

```
#define BARRY_SAFE_EXP -100.0
```

Definition at line 33 of file barry-configuration.hpp.

8.18.1.5 printf_barry

```
#define printf_barry printf
```

Definition at line 51 of file barry-configuration.hpp.

8.18.2 Typedef Documentation

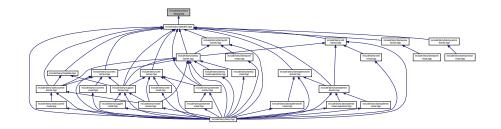
8.18.2.1 Map

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

Definition at line 27 of file barry-configuration.hpp.

8.19 include/barry/barry-debug.hpp File Reference

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



Macros

• #define BARRY_DEBUG_LEVEL 0

8.19.1 Macro Definition Documentation

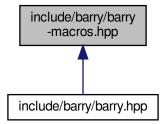
8.19.1.1 BARRY_DEBUG_LEVEL

#define BARRY_DEBUG_LEVEL 0

Definition at line 5 of file barry-debug.hpp.

8.20 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);

8.20.1 Macro Definition Documentation

8.20.1.1 BARRY_ONE

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

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

8.20.1.2 BARRY_ONE_DENSE

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

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

8.20.1.3 BARRY_UNUSED

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

8.20.1.4 BARRY_ZERO

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

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

8.20.1.5 BARRY_ZERO_DENSE

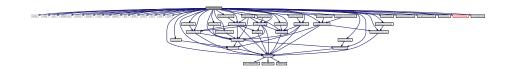
```
#define BARRY_ZERO_DENSE static_cast<Cell_Type>(0.0)
```

Definition at line 5 of file barry-macros.hpp.

8.21 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 "typedefs.hpp"
#include "barry-macros.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 "counters/phylo.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
- barry::counters::phylo

Macros

- #define BARRY HPP
- #define BARRY VERSION 0.1
- #define COUNTER_FUNCTION(a)
- #define COUNTER_LAMBDA(a)
- #define RULE_FUNCTION(a)
- #define RULE_LAMBDA(a)

8.21.1 Macro Definition Documentation

8.21.1.1 BARRY_HPP

#define BARRY_HPP

Definition at line 22 of file barry.hpp.

8.21.1.2 BARRY_VERSION

#define BARRY_VERSION 0.1

Definition at line 24 of file barry.hpp.

8.21.1.3 COUNTER_FUNCTION

Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
inline double (a) (const Array_Type & Array, uint i, uint j, Data_Type & data) \
```

Definition at line 86 of file barry.hpp.

8.21.1.4 COUNTER_LAMBDA

Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
Counter_fun_type<Array_Type, Data_Type> a = \
[](const Array_Type & Array, uint i, uint j, Data_Type & data)
```

Definition at line 89 of file barry.hpp.

8.21.1.5 RULE FUNCTION

Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
inline bool (a) (const Array_Type & Array, uint i, uint j, Data_Type & data) \
```

Definition at line 93 of file barry.hpp.

8.21.1.6 **RULE LAMBDA**

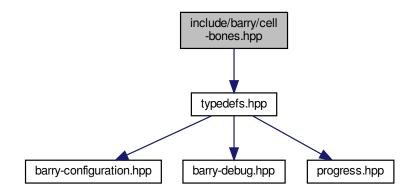
Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
Rule_fun_type<Array_Type, Data_Type> a = \
[](const Array_Type & Array, uint i, uint j, Data_Type & data)
```

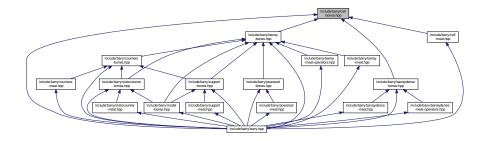
Definition at line 96 of file barry.hpp.

8.22 include/barry/cell-bones.hpp File Reference

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



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



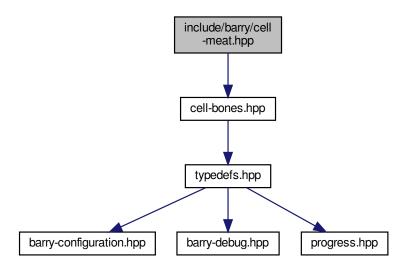
Classes

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

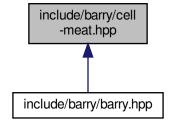
8.23 include/barry/cell-meat.hpp File Reference

#include "cell-bones.hpp"

Include dependency graph for cell-meat.hpp:



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



8.24 include/barry/col-bones.hpp File Reference

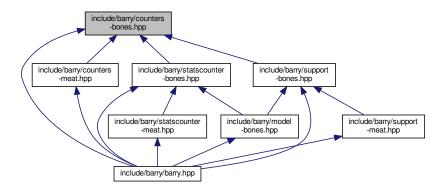
8.25 include/barry/counters-bones.hpp File Reference

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

Include dependency graph for counters-bones.hpp:



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

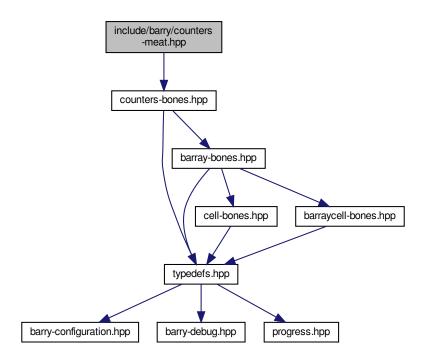


Classes

- class Counters
 Array_Type, Data_Type >
 Vector of counters.

8.26 include/barry/counters-meat.hpp File Reference

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



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



Macros

- #define COUNTER_TYPE() Counter<Array_Type,Data_Type>
- #define COUNTER_TEMPLATE_ARGS() < typename Array_Type, typename Data_Type>

- #define COUNTER_TEMPLATE(a, b) template COUNTER_TEMPLATE_ARGS() inline a COUNTER_TYPE() ↔ ::b
- #define COUNTERS_TYPE() Counters<Array_Type,Data_Type>
- #define COUNTERS_TEMPLATE_ARGS() < typename Array_Type, typename Data_Type>

Functions

- COUNTER_TEMPLATE (, Counter)(const Counter< Array_Type
- Data Type init fun (counter .init fun)
- Data Type &&counter init fun (std::move(counter .init 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
- COUNTERS_TEMPLATE (, Counters)()
- COUNTERS_TEMPLATE (COUNTER_TYPE() &, operator[])(uint idx)
- Data_Type Counters (Counters < Array_Type, Data_Type > &&counters_) noexcept
- 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

Variables

- Data Type & counter
- Data_Type &&counter_ noexcept
- uint i
- · uint uint j
- return * this
- 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 > Data_Type data_
- Data_Type Counter_fun_type
 Array_Type, Data_Type > Data_Type std::string name_
- Data_Type Counter_fun_type
 Array_Type, Data_Type > Data_Type std::string std::string desc_

8.26.1 Macro Definition Documentation

8.26.1.1 COUNTER_TEMPLATE

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

8.26.1.2 COUNTER_TEMPLATE_ARGS

```
#define COUNTER_TEMPLATE_ARGS() <typename Array_Type, typename Data_Type>
```

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

8.26.1.3 COUNTER_TYPE

```
#define COUNTER_TYPE( ) Counter<Array_Type, Data_Type>
```

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

8.26.1.4 COUNTERS_TEMPLATE

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

8.26.1.5 COUNTERS_TEMPLATE_ARGS

```
#define COUNTERS_TEMPLATE_ARGS() <typename Array_Type, typename Data_Type>
```

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

8.26.1.6 COUNTERS_TYPE

```
#define COUNTERS_TYPE( ) Counters<Array_Type,Data_Type>
```

Definition at line 114 of file counters-meat.hpp.

8.26.2 Function Documentation

8.26.2.1 count fun()

8.26.2.2 COUNTER_TEMPLATE() [1/7]

```
COUNTER_TEMPLATE (

Counter ) const
```

8.26.2.3 COUNTER_TEMPLATE() [2/7]

8.26.2.4 COUNTER_TEMPLATE() [3/7]

8.26.2.5 COUNTER_TEMPLATE() [4/7]

< Move assignment

8.26.2.6 COUNTER_TEMPLATE() [5/7]

```
COUNTER_TEMPLATE ( \label{eq:counter_template} \mbox{double ,} \\ \mbox{init ) & } \&
```

8.26.2.7 COUNTER_TEMPLATE() [6/7]

```
COUNTER_TEMPLATE (
          std::string ,
          get_description ) const
```

Definition at line 106 of file counters-meat.hpp.

8.26.2.8 COUNTER_TEMPLATE() [7/7]

```
COUNTER_TEMPLATE (
          std::string ,
          get_name ) const
```

Definition at line 102 of file counters-meat.hpp.

8.26.2.9 Counters()

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

8.26.2.10 COUNTERS_TEMPLATE() [1/7]

```
COUNTERS_TEMPLATE (

Counters )
```

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

8.26.2.11 COUNTERS_TEMPLATE() [2/7]

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

8.26.2.12 COUNTERS_TEMPLATE() [3/7]

```
COUNTERS_TEMPLATE (

COUNTERS_TYPE() & ,

operator )
```

8.26.2.13 COUNTERS_TEMPLATE() [4/7]

```
COUNTERS_TEMPLATE (

COUNTERS_TYPE() ,

operator ) const
```

8.26.2.14 COUNTERS_TEMPLATE() [5/7]

```
COUNTERS_TEMPLATE (
          std::vector< std::string > ,
          get_descriptions ) const
```

Definition at line 194 of file counters-meat.hpp.

8.26.2.15 COUNTERS_TEMPLATE() [6/7]

```
COUNTERS_TEMPLATE (
          std::vector< std::string > ,
          get_names ) const
```

Definition at line 183 of file counters-meat.hpp.

8.26.2.16 COUNTERS_TEMPLATE() [7/7]

```
COUNTERS_TEMPLATE (
     void ,
     add_counter )
```

8.26.2.17 data()

8.26.2.18 desc()

Move constructor.

Definition at line 33 of file counters-meat.hpp.

8.26.2.19 init_fun() [1/3]

8.26.2.20 init_fun() [2/3]

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

8.26.2.21 init_fun() [3/3]

8.26.2.22 name()

8.26.3 Variable Documentation

8.26.3.1 count_fun_

```
Data_Type count_fun_
```

Definition at line 163 of file counters-meat.hpp.

8.26.3.2 counter

```
Data_Type counter

Initial value:
{
    data.push_back(counter)
```

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

8.26.3.3 counter_

```
Data_Type & counter_

Initial value:
{
    if (this != &counter_) {
        this->count_fun = counter_.count_fun;
        this->init_fun = counter_.init_fun;

        this->data = counter_.data;
        this->name = counter_.name;
        this->desc = counter_.desc;
    }
    return *this
```

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

8.26.3.4 data_

```
Data_Type Counter_fun_type<Array_Type,Data_Type> Data_Type data_
```

Definition at line 165 of file counters-meat.hpp.

8.26.3.5 desc

Data_Type Counter_fun_type<Array_Type,Data_Type > Data_Type std::string std::string desc_

Initial value:

```
{
```

```
data.push_back(Counter<Array_Type,Data_Type>(
    count_fun_,
    init_fun_,
    data_,
    name_,
    desc_
```

Definition at line 167 of file counters-meat.hpp.

8.26.3.6 i

uint i

Definition at line 82 of file counters-meat.hpp.

8.26.3.7 init_fun_

```
Data_Type Counter_fun_type<Array_Type,Data_Type> init_fun_
```

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

8.26.3.8 j

```
uint uint j
```

Initial value:

```
if (count_fun == nullptr)
    return 0.0
```

Definition at line 82 of file counters-meat.hpp.

8.26.3.9 name_

```
Data_Type Counter_fun_type<Array_Type,Data_Type> Data_Type std::string name_
```

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

8.26.3.10 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->name = std::move(counter_.name);
    this->desc = std::move(counter_.desc);
}
return *this
```

Definition at line 28 of file counters-meat.hpp.

8.26.3.11 return

return

Definition at line 159 of file counters-meat.hpp.

8.26.3.12 this

```
return* this
```

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

8.27 include/barry/counters/defm.hpp File Reference

Classes

• class DEFMData

Data class for DEFM arrays.

· class DEFMCounterData

Data class used to store arbitrary uint or double vectors.

Macros

Macros for defining counters

- #define DEFM COUNTER(a)
- #define DEFM COUNTER LAMBDA(a)
- #define NETWORKDENSE_COUNTER_LAMBDA(a)

Macros for defining rules

- #define DEFM RULE(a)
- #define DEFM_RULE_LAMBDA(a)

Typedefs

Convenient typedefs for network objects.

```
    typedef BArrayDense < double, DEFMData > DEFMArray
    template < typename Tarray = DEFMArray > using DEFMCounter = Counter < Tarray, DEFMCounterData >
    template < typename Tarray = DEFMArray > using DEFMCounters = Counters < Tarray, DEFMCounterData >
    template < typename Tarray = DEFMArray > using DEFMSupport = Support < Tarray, DEFMCounterData >
    template < typename Tarray = DEFMArray > using DEFMStatsCounter = StatsCounter < Tarray, DEFMCounterData >
    template < typename Tarray > using DEFMModel = Model < Tarray, DEFMCounterData >
    template < typename Tarray = DEFMArray > using DEFMRule = Rule < Tarray, bool >
    template < typename Tarray = DEFMArray > using DEFMRule = Rule < Tarray, bool >
```

Functions

```
    template < typename Tarray = DEFMArray > void counter_ones (DEFMCounters < Tarray > *counters, int covar_index=-1)
    Prevalence of ones.
```

Rules for network models

Parameters

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

```
    template<typename Tarray = DEFMArray>
    void rules_zerodiag (DEFMRules< Tarray > *rules)
    Number of edges.
```

8.27.1 Macro Definition Documentation

8.27.1.1 DEFM_COUNTER

Function for definition of a network counter function

Definition at line 123 of file defm.hpp.

8.27.1.2 DEFM COUNTER LAMBDA

Lambda function for definition of a network counter function

Definition at line 128 of file defm.hpp.

8.27.1.3 DEFM RULE

Function for definition of a network counter function

Definition at line 142 of file defm.hpp.

8.27.1.4 DEFM_RULE_LAMBDA

Lambda function for definition of a network counter function

Definition at line 147 of file defm.hpp.

8.27.1.5 NETWORKDENSE_COUNTER_LAMBDA

Definition at line 132 of file defm.hpp.

8.27.2 Typedef Documentation

8.27.2.1 DEFMArray

```
typedef BArrayDense<double, DEFMData> DEFMArray
```

Definition at line 95 of file defm.hpp.

8.27.2.2 DEFMCounter

```
template<typename Tarray = DEFMArray>
using DEFMCounter = Counter<Tarray, DEFMCounterData >
```

Definition at line 98 of file defm.hpp.

8.27.2.3 DEFMCounters

```
template<typename Tarray = DEFMArray>
using DEFMCounters = Counters<Tarray, DEFMCounterData>
```

Definition at line 101 of file defm.hpp.

8.27.2.4 DEFMModel

```
template<typename Tarray >
using DEFMModel = Model<Tarray, DEFMCounterData>
```

Definition at line 110 of file defm.hpp.

8.27.2.5 **DEFMRule**

```
template<typename Tarray = DEFMArray>
using DEFMRule = Rule<Tarray, bool>
```

Definition at line 113 of file defm.hpp.

8.27.2.6 **DEFMRules**

```
template<typename Tarray = DEFMArray>
using DEFMRules = Rules<Tarray, bool>
```

Definition at line 116 of file defm.hpp.

8.27.2.7 DEFMStatsCounter

```
template<typename Tarray = DEFMArray>
using DEFMStatsCounter = StatsCounter<Tarray, DEFMCounterData>
```

Definition at line 107 of file defm.hpp.

8.27.2.8 DEFMSupport

```
template<typename Tarray = DEFMArray>
using DEFMSupport = Support<Tarray, DEFMCounterData >
```

Definition at line 104 of file defm.hpp.

8.27.3 Function Documentation

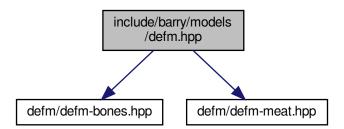
8.27.3.1 rules_zerodiag()

Number of edges.

Definition at line 258 of file defm.hpp.

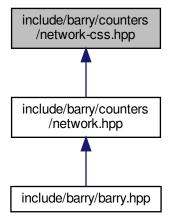
8.28 include/barry/models/defm.hpp File Reference

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



8.29 include/barry/counters/network-css.hpp File Reference

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



Macros

- #define CSS_SIZE()
- #define CSS_CASE_TRUTH() if ((i < n) && (j < n))
- #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, uint netsize, const std↔
  ::vector< uint > &end_)
     Counts errors of commission.
• template<typename Tnet = Network>
  void counter css partially false recip omiss (NetCounters< Tnet > *counters, uint netsize, const std↔
  ::vector< uint > &end )
     Counts errors of omission.
• template<typename Tnet = Network>
  void counter_css_completely_false_recip_comiss (NetCounters< Tnet > *counters, uint netsize, const std ←
  ::vector< uint > &end )
     Counts completely false reciprocity (comission)
• template<typename Tnet = Network>
  void counter_css_completely_false_recip_omiss (NetCounters< Tnet > *counters, uint netsize, const std↔
  ::vector< uint > &end_)
     Counts completely false reciprocity (omission)

    template<typename Tnet = Network>

  void counter_css_mixed_recip (NetCounters< Tnet > *counters, uint netsize, const std::vector< uint >
  &end_)
     Counts mixed reciprocity errors.
• template<typename Tnet = Network>
  void counter css census01 (NetCounters< Tnet > *counters, uint netsize, const std::vector< uint > &end ←
template<typename Tnet = Network>
  void counter css census02 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
  _)
• template<typename Tnet = Network>
 void counter css census03 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
template<typename Tnet = Network>
  void counter_css_census04 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
 _)

    template<tvpename Tnet = Network>

 void counter_css_census05 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
template<typename Tnet = Network>
 void counter css census06 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
template<typename Tnet = Network>
  void counter_css_census07 (NetCounters< Tnet > *counters, uint netsize, const std::vector< uint > &end↔
 _)
```

void counter_css_census08 (NetCounters< Tnet > *counters, uint netsize, const std::vector< uint > &end↔

_)

template<typename Tnet = Network>

```
    template<typename Tnet = Network>
        void counter_css_census09 (NetCounters< Tnet > *counters, uint netsize, const std::vector< uint > &end
        —)
```

• template<typename Tnet = Network> void counter_css_census10 (NetCounters< Tnet > *counters, uint netsize, const std::vector< uint > &end ← _)

8.29.1 Macro Definition Documentation

8.29.1.1 CSS_APPEND

Value:

Definition at line 42 of file network-css.hpp.

8.29.1.2 CSS_CASE_ELSE

```
#define CSS_CASE_ELSE( )
```

Definition at line 27 of file network-css.hpp.

8.29.1.3 CSS CASE PERCEIVED

```
#define CSS_CASE_PERCEIVED() else if (((i >= s) && (i < e)) & ((j >= s) && (j < e)))
```

Definition at line 20 of file network-css.hpp.

8.29.1.4 CSS CASE TRUTH

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

Definition at line 13 of file network-css.hpp.

8.29.1.5 CSS_CHECK_SIZE

Definition at line 37 of file network-css.hpp.

8.29.1.6 CSS_CHECK_SIZE_INIT

Definition at line 31 of file network-css.hpp.

8.29.1.7 CSS_NET_COUNTER_LAMBDA_INIT

Definition at line 49 of file network-css.hpp.

8.29.1.8 CSS PERCEIVED CELLS

```
#define CSS_PERCEIVED_CELLS()

Value:
    double tji = static_cast<double>(Array(j - s, i - s, false)); \
    double pji = static_cast<double>(Array(j, i, false)); \
    double tij = static_cast<double>(Array(i - s, j - s, false));
```

Definition at line 21 of file network-css.hpp.

8.29.1.9 CSS_SIZE

```
#define CSS_SIZE( )

Value:
    uint n = data.indices[0u]; \
    uint s = data.indices[1u]; \
    uint e = data.indices[2u];
```

Definition at line 7 of file network-css.hpp.

8.29.1.10 CSS TRUE CELLS

```
#define CSS_TRUE_CELLS( )

Value:
    double tji = static_cast<double>(Array(j, i, false)); \
    double pij = static_cast<double>(Array(i + s, j + s, false)); \
    double pji = static_cast<double>(Array(j + s, i + s, false));
```

Definition at line 14 of file network-css.hpp.

8.29.2 Function Documentation

8.29.2.1 counter css census01()

Definition at line 275 of file network-css.hpp.

8.29.2.2 counter_css_census02()

Definition at line 325 of file network-css.hpp.

8.29.2.3 counter_css_census03()

Definition at line 364 of file network-css.hpp.

8.29.2.4 counter_css_census04()

Definition at line 403 of file network-css.hpp.

8.29.2.5 counter_css_census05()

Definition at line 442 of file network-css.hpp.

8.29.2.6 counter_css_census06()

Definition at line 481 of file network-css.hpp.

8.29.2.7 counter_css_census07()

Definition at line 520 of file network-css.hpp.

8.29.2.8 counter_css_census08()

Definition at line 559 of file network-css.hpp.

8.29.2.9 counter_css_census09()

Definition at line 598 of file network-css.hpp.

8.29.2.10 counter_css_census10()

Definition at line 637 of file network-css.hpp.

8.29.2.11 counter_css_completely_false_recip_comiss()

Counts completely false reciprocity (comission)

Definition at line 154 of file network-css.hpp.

8.29.2.12 counter_css_completely_false_recip_omiss()

Counts completely false reciprocity (omission)

Definition at line 194 of file network-css.hpp.

8.29.2.13 counter css mixed recip()

Counts mixed reciprocity errors.

Definition at line 234 of file network-css.hpp.

8.29.2.14 counter_css_partially_false_recip_commi()

Counts errors of commission.

Parameters

netsize	Size of the reference (true) network
end⊷	Vector indicating one past the ending index of each network. (see details)
_	

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

Definition at line 63 of file network-css.hpp.

8.29.2.15 counter_css_partially_false_recip_omiss()

Counts errors of omission.

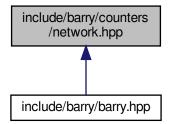
Definition at line 110 of file network-css.hpp.

8.30 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 uint or double vectors.

Macros

- #define NET_C_DATA_IDX(i) (data.indices[i])
- #define NET_C_DATA_NUM(i) (data.numbers[i])

Macros for defining counters

- #define NETWORK_COUNTER(a)
- #define NETWORK_COUNTER_LAMBDA(a)
- #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, uint 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, uint attr id, double alpha=1.0, double tail head=true)
     Sum of attribute difference between ego and alter to pow(alpha)

    NETWORK_COUNTER (init_single_attr)

    template<typename Tnet = Network>

  void counter nodeicov (NetCounters < Tnet > *counters, uint attr id)
template<typename Tnet = Network>
  void counter nodeocov (NetCounters< Tnet > *counters, uint attr id)
template<typename Tnet = Network>
  void counter_nodecov (NetCounters< Tnet > *counters, uint attr_id)
template<typename Tnet = Network>
  void counter_nodematch (NetCounters < Tnet > *counters, uint attr_id)
• template<typename Tnet = Network>
  void counter_idegree (NetCounters< Tnet > *counters, std::vector< uint > d)
     Counts number of vertices with a given in-degree.

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

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

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

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

Rules for network models

Parameters

rules | A pointer to a NetRules object (Rules < Network, bool >).

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 >

8.30.1 Macro Definition Documentation

8.30.1.1 BARRY_ZERO_NETWORK

#define BARRY ZERO NETWORK 0.0

Definition at line 85 of file network.hpp.

8.30.1.2 BARRY_ZERO_NETWORK_DENSE

```
#define BARRY_ZERO_NETWORK_DENSE 0
```

Definition at line 86 of file network.hpp.

8.30.1.3 NET_C_DATA_IDX

Definition at line 74 of file network.hpp.

8.30.1.4 NET_C_DATA_NUM

Definition at line 75 of file network.hpp.

8.30.1.5 NETWORK_COUNTER

Value:

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

Function for definition of a network counter function

Definition at line 114 of file network.hpp.

8.30.1.6 NETWORK_COUNTER_LAMBDA

Value:

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

Lambda function for definition of a network counter function

Definition at line 119 of file network.hpp.

8.30.1.7 NETWORK_RULE

inline bool (a) (const Tnet & Array, uint i, uint j, bool & data)

Function for definition of a network counter function

Definition at line 133 of file network.hpp.

8.30.1.8 NETWORK RULE LAMBDA

Lambda function for definition of a network counter function

Definition at line 138 of file network.hpp.

8.30.1.9 NETWORKDENSE_COUNTER_LAMBDA

Definition at line 123 of file network.hpp.

8.30.2 Typedef Documentation

8.30.2.1 NetCounter

```
template<typename Tnet = Network>
using NetCounter = Counter<Tnet, NetCounterData >
```

Definition at line 89 of file network.hpp.

8.30.2.2 NetCounters

```
template<typename Tnet = Network>
using NetCounters = Counters<Tnet, NetCounterData>
```

Definition at line 92 of file network.hpp.

8.30.2.3 NetModel

```
template<typename Tnet >
using NetModel = Model<Tnet, NetCounterData>
```

Definition at line 101 of file network.hpp.

8.30.2.4 NetRule

```
template<typename Tnet = Network>
using NetRule = Rule<Tnet, bool>
```

Definition at line 104 of file network.hpp.

8.30.2.5 NetRules

```
template<typename Tnet = Network>
using NetRules = Rules<Tnet, bool>
```

Definition at line 107 of file network.hpp.

8.30.2.6 NetStatsCounter

```
template<typename Tnet = Network>
using NetStatsCounter = StatsCounter<Tnet, NetCounterData>
```

Definition at line 98 of file network.hpp.

8.30.2.7 NetSupport

```
template<typename Tnet = Network>
using NetSupport = Support<Tnet, NetCounterData >
```

Definition at line 95 of file network.hpp.

8.30.2.8 Network

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

Definition at line 82 of file network.hpp.

8.30.2.9 NetworkDense

```
typedef BArrayDense<int, NetworkData> NetworkDense
```

Definition at line 83 of file network.hpp.

8.30.3 Function Documentation

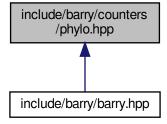
8.30.3.1 rules_zerodiag()

Number of edges.

Definition at line 1383 of file network.hpp.

8.31 include/barry/counters/phylo.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 DEFAULT_DUPLICATION 1u
- #define DUPL SPEC 0u
- #define DUPL_DUPL 1u
- #define DUPL EITH 2u
- #define MAKE DUPL VARS()
- #define IS EITHER() (DATA AT == DUPL EITH)
- #define IS_DUPLICATION() ((DATA_AT == DUPL_DUPL) & (DPL))
- #define IS_SPECIATION() ((DATA_AT == DUPL_SPEC) & (!DPL))
- #define IF MATCHES()
- #define IF_NOTMATCHES()
- #define PHYLO COUNTER LAMBDA(a)

Extension of a simple counter.

- #define PHYLO_RULE_DYN_LAMBDA(a)
- #define PHYLO_CHECK_MISSING()

Typedefs

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

Convenient typedefs for Node objects.

- typedef BArrayDense< uint, NodeData > PhyloArray
- typedef Counter
 PhyloArray, PhyloCounterData > PhyloCounter
- typedef Counters< PhyloArray, PhyloCounterData > PhyloCounters
- typedef Rule < PhyloArray, PhyloRuleData > PhyloRule
- typedef Rules
 PhyloArray, PhyloRuleData > PhyloRules
- typedef Rule
 PhyloArray, PhyloRuleDynData
 PhyloRuleDyn
- typedef Rules
 PhyloArray, PhyloRuleDynData
 PhyloRulesDyn
- typedef Support < PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloSupport
- typedef StatsCounter< PhyloArray, PhyloCounterData > PhyloStatsCounter
- typedef Model < PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloModel
- typedef PowerSet < PhyloArray, PhyloRuleData > PhyloPowerSet

Functions

- std::string get last name (unsigned int d)
- void counter_overall_gains (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)
 Overall functional gains.
- void counter_gains (PhyloCounters *counters, std::vector< uint > nfun, unsigned int duplication=DEFAULT_DUPLICATION)
 Functional gains for a specific function (nfun).
- void counter_gains_k_offspring (PhyloCounters *counters, std::vector< uint > nfun, uint k=1u, unsigned int duplication=DEFAULT DUPLICATION)

k genes gain function nfun

- void counter_genes_changing (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)
- void counter_preserve_pseudogene (PhyloCounters *counters, unsigned int nfunA, unsigned int nfunB, unsigned int duplication=DEFAULT DUPLICATION)

Keeps track of how many pairs of genes preserve pseudostate.

- void counter_prop_genes_changing (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)
- void counter_overall_loss (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)
 Overall functional loss.
- void counter_maxfuns (PhyloCounters *counters, uint lb, uint ub, unsigned int duplication=DEFAULT_DUPLICATION)

 Cap the number of functions per gene.
- void counter_loss (PhyloCounters *counters, std::vector < uint > nfun, unsigned int duplication=DEFAULT_DUPLICATION)

 Total count of losses for an specific function.
- void counter_overall_changes (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Total number of changes. Use this statistic to account for "preservation".
- void counter_subfun (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

 Total count of Sub-functionalization events.
- void counter_cogain (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

 Co-evolution (joint gain or loss)
- void counter_longest (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)
 Longest branch mutates (either by gain or by loss)
- void counter_neofun (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

 Total number of neofunctionalization events.
- void counter_pairwise_neofun_singlefun (PhyloCounters *counters, uint nfunA, unsigned int duplication=DEFAULT_DUPLICATION 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, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION Total number of neofunctionalization events.
- void counter_co_opt (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION) Function co-opting.
- void counter_k_genes_changing (PhyloCounters *counters, unsigned int k, unsigned int duplication=DEFAULT_DUPLICATION)

 Indicator function. Equals to one if k genes changed and zero otherwise.
- void counter_less_than_p_prop_genes_changing (PhyloCounters *counters, double p, unsigned int duplication=DEFAULT DUPLICATION)

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

- void counter_gains_from_0 (PhyloCounters *counters, std::vector < uint > nfun, unsigned int duplication=DEFAULT_DUPLICAT
 Used when all the functions are in 0 (like the root node prob.)
- void counter_overall_gains_from_0 (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Used when all the functions are in 0 (like the root node prob.)
- void counter_pairwise_overall_change (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

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

 void counter_pairwise_preserving (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

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

 void counter_pairwise_first_gain (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

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

void rule_dyn_limit_changes (PhyloSupport *support, uint pos, uint lb, uint ub, unsigned int duplication=DEFAULT_DUPLICATIC
 Overall functional gains.

8.31.1 Macro Definition Documentation

8.31.1.1 DEFAULT_DUPLICATION

#define DEFAULT_DUPLICATION 1u

Definition at line 5 of file phylo.hpp.

8.31.1.2 DUPL DUPL

#define DUPL_DUPL 1u

Definition at line 7 of file phylo.hpp.

8.31.1.3 DUPL_EITH

#define DUPL_EITH 2u

Definition at line 8 of file phylo.hpp.

8.31.1.4 DUPL_SPEC

#define DUPL_SPEC Ou

Definition at line 6 of file phylo.hpp.

8.31.1.5 IF_MATCHES

```
#define IF_MATCHES( )

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

Definition at line 19 of file phylo.hpp.

8.31.1.6 IF_NOTMATCHES

```
#define IF_NOTMATCHES( )

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

Definition at line 21 of file phylo.hpp.

8.31.1.7 IS_DUPLICATION

```
#define IS_DUPLICATION( ) ((DATA_AT == DUPL_DUPL) & (DPL))
```

Definition at line 16 of file phylo.hpp.

8.31.1.8 IS_EITHER

```
#define IS_EITHER( ) (DATA_AT == DUPL_EITH)
```

Definition at line 15 of file phylo.hpp.

8.31.1.9 IS_SPECIATION

```
#define IS_SPECIATION( ) ((DATA_AT == DUPL_SPEC) & (!DPL))
```

Definition at line 17 of file phylo.hpp.

8.31.1.10 MAKE_DUPL_VARS

```
#define MAKE_DUPL_VARS( )

Value:
   bool DPL = Array.D()->duplication; \
   unsigned int DATA_AT = data[Ou];
```

Definition at line 11 of file phylo.hpp.

8.31.1.11 PHYLO_CHECK_MISSING

```
#define PHYLO_CHECK_MISSING( )

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

Definition at line 139 of file phylo.hpp.

8.31.1.12 PHYLO_COUNTER_LAMBDA

Extension of a simple counter.

It allows specifying extra arguments, in particular, the corresponding sets of rows to which this statistic may be relevant. This could be important in the case of, for example, counting correlation type statistics between function 1 and 2, and between function 1 and 3.

Definition at line 133 of file phylo.hpp.

8.31.1.13 PHYLO_RULE_DYN_LAMBDA

Definition at line 136 of file phylo.hpp.

8.31.2 Typedef Documentation

8.31.2.1 PhyloArray

typedef BArrayDense<uint, NodeData> PhyloArray

Definition at line 106 of file phylo.hpp.

8.31.2.2 PhyloCounter

typedef Counter<PhyloArray, PhyloCounterData > PhyloCounter

Definition at line 107 of file phylo.hpp.

8.31.2.3 PhyloCounters

typedef Counters< PhyloArray, PhyloCounterData> PhyloCounters

Definition at line 108 of file phylo.hpp.

8.31.2.4 PhyloModel

 ${\tt typedef\ Model < PhyloArray,\ PhyloCounterData,\ PhyloRuleData,\ PhyloRuleDynData > PhyloModel}$

Definition at line 118 of file phylo.hpp.

8.31.2.5 PhyloPowerSet

typedef PowerSet<PhyloArray, PhyloRuleData> PhyloPowerSet

Definition at line 119 of file phylo.hpp.

8.31.2.6 PhyloRule

typedef Rule<PhyloArray,PhyloRuleData> PhyloRule

Definition at line 110 of file phylo.hpp.

8.31.2.7 PhyloRuleData

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

Definition at line 99 of file phylo.hpp.

8.31.2.8 PhyloRuleDyn

typedef Rule<PhyloArray,PhyloRuleDynData> PhyloRuleDyn

Definition at line 113 of file phylo.hpp.

8.31.2.9 PhyloRules

typedef Rules<PhyloArray,PhyloRuleData> PhyloRules

Definition at line 111 of file phylo.hpp.

8.31.2.10 PhyloRulesDyn

typedef Rules<PhyloArray,PhyloRuleDynData> PhyloRulesDyn

Definition at line 114 of file phylo.hpp.

8.31.2.11 PhyloStatsCounter

typedef StatsCounter<PhyloArray, PhyloCounterData> PhyloStatsCounter

Definition at line 117 of file phylo.hpp.

8.31.2.12 PhyloSupport

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

Definition at line 116 of file phylo.hpp.

8.31.3 Function Documentation

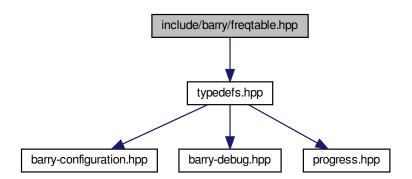
8.31.3.1 get_last_name()

```
std::string get_last_name (
          unsigned int d ) [inline]
```

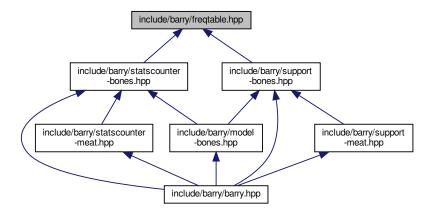
Definition at line 142 of file phylo.hpp.

8.32 include/barry/freqtable.hpp File Reference

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



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



Classes

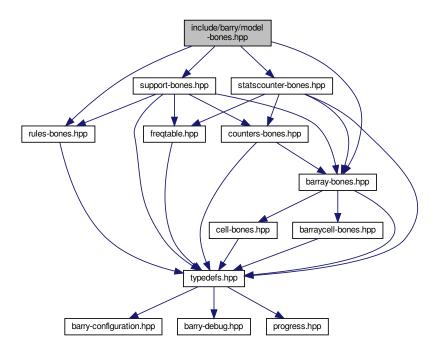
class FreqTable
 T >

Frequency table of vectors.

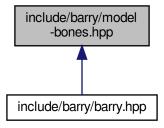
8.33 include/barry/model-bones.hpp File Reference

```
#include "barray-bones.hpp"
#include "support-bones.hpp"
#include "statscounter-bones.hpp"
#include "rules-bones.hpp"
```

Include dependency graph for model-bones.hpp:



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



Classes

class Model < Array_Type, Data_Counter_Type, Data_Rule_Type, 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:

Functions

template<typename Array_Type >
 std::vector< double > keygen_default (const Array_Type &Array_)
 Array Hasher class (used for computing support)

8.33.1 Function Documentation

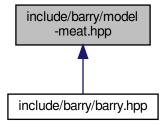
8.33.1.1 keygen_default()

Array Hasher class (used for computing support)

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

8.34 include/barry/model-meat.hpp File Reference

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



Macros

- #define MODEL_TYPE()
- #define MODEL TEMPLATE ARGS()
- #define MODEL_TEMPLATE(a, b) template MODEL_TEMPLATE_ARGS() inline a MODEL_TYPE()::b

Functions

- double update_normalizing_constant (const double *params, const double *support, size_t k, size_t n)
- double likelihood_ (const double *stats_target, const std::vector< double > ¶ms, const double normalizing_constant, size_t n_params, bool log_=false)
- MODEL_TEMPLATE (, Model)()
- MODEL_TEMPLATE (, Model)(const MODEL_TYPE() &Model_)

8.34.1 Macro Definition Documentation

8.34.1.1 MODEL_TEMPLATE

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

8.34.1.2 MODEL_TEMPLATE_ARGS

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

8.34.1.3 MODEL_TYPE

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

8.34.2 Function Documentation

8.34.2.1 likelihood_()

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

8.34.2.2 MODEL_TEMPLATE() [1/2]

```
MODEL_TEMPLATE (

Model )
```

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

8.34.2.3 MODEL_TEMPLATE() [2/2]

```
MODEL_TEMPLATE (

Model ) const &
```

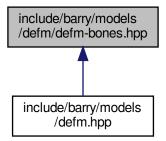
Definition at line 157 of file model-meat.hpp.

8.34.2.4 update_normalizing_constant()

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

8.35 include/barry/models/defm/defm-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

- RULE_FUNCTION (rule_empty_free)
- std::vector< double > keygen_full (const phylocounters::PhyloArray &array)
- bool vec_diff (const std::vector< unsigned int > &s, const std::vector< unsigned int > &a)

8.35.1 Macro Definition Documentation

8.35.1.1 INITIALIZED

```
#define INITIALIZED( )

Value:
    if (!this->initialized) \
        throw std::logic_error("The model has not been initialized yet.");
```

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

8.35.2 Function Documentation

8.35.2.1 keygen_full()

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

8.35.2.2 RULE_FUNCTION()

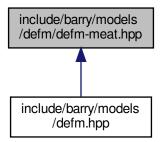
Definition at line 13 of file defm-bones.hpp.

8.35.2.3 vec_diff()

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

8.36 include/barry/models/defm/defm-meat.hpp File Reference

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



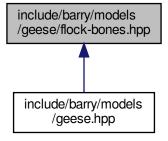
8.37 include/barry/models/geese.hpp File Reference

```
#include "geese/geese-node-bones.hpp"
#include "geese/geese-bones.hpp"
#include "geese/geese-meat.hpp"
#include "geese/geese-meat-constructors.hpp"
#include "geese/geese-meat-likelihood.hpp"
#include "geese/geese-meat-likelihood_exhaust.hpp"
#include "geese/geese-meat-simulate.hpp"
#include "geese/geese-meat-predict.hpp"
#include "geese/geese-meat-predict_exhaust.hpp"
#include "geese/geese-meat-predict_sim.hpp"
#include "geese/flock-bones.hpp"
#include "geese/flock-meat.hpp"
Include dependency graph for geese.hpp:
```



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

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

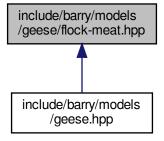


Classes

class Flock

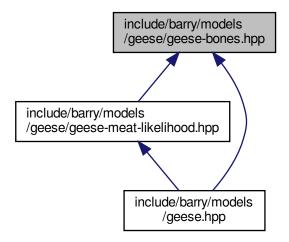
A Flock is a group of Geese.

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



8.40 include/barry/models/geese/geese-bones.hpp File Reference

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



Classes

class Geese

Annotated Phylo Model.

Macros

• #define INITIALIZED()

Functions

- template<typename Ta , typename Tb > $std::vector < Ta > vector_caster \ (const \ std::vector < Tb > \&x)$
- RULE_FUNCTION (rule_empty_free)
- std::vector< double > keygen_full (const phylocounters::PhyloArray &array)
- $\bullet \ \ \mathsf{bool} \ \mathsf{vec_diff} \ (\mathsf{const} \ \mathsf{std} : \mathsf{vector} < \mathsf{unsigned} \ \mathsf{int} > \mathsf{\&s}, \ \mathsf{const} \ \mathsf{std} : \mathsf{vector} < \mathsf{unsigned} \ \mathsf{int} > \mathsf{\&a}) \\$

8.40.1 Macro Definition Documentation

8.40.1.1 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.

8.40.2 Function Documentation

8.40.2.1 keygen_full()

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

8.40.2.2 RULE_FUNCTION()

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

8.40.2.3 vec_diff()

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

8.40.2.4 vector_caster()

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

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

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

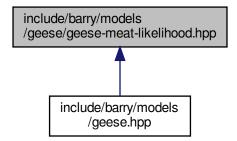


8.42 include/barry/models/geese/geese-meat-likelihood.hpp File Reference

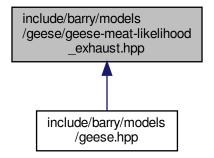
#include "geese-bones.hpp"
Include dependency graph for geese-meat-likelihood.hpp:



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



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



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

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

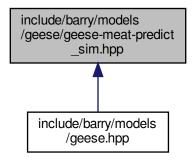


8.45 include/barry/models/geese/geese-meat-predict_exhaust.hpp File Reference



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

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

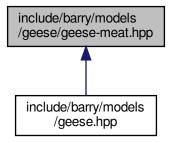


8.47 include/barry/models/geese/geese-meat-simulate.hpp File Reference



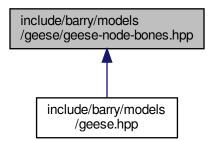
8.48 include/barry/models/geese/geese-meat.hpp File Reference

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



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

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



Classes

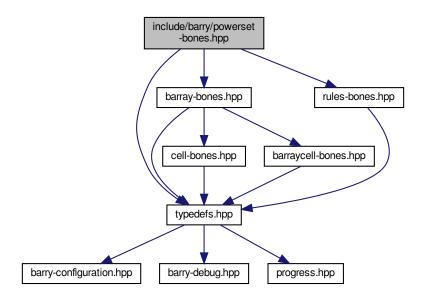
• class Node

A single node for the model.

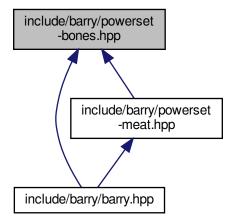
8.50 include/barry/powerset-bones.hpp File Reference

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

Include dependency graph for powerset-bones.hpp:



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



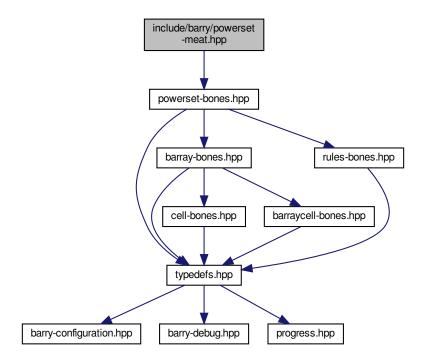
Classes

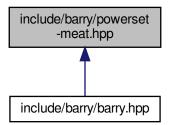
class PowerSet
 Array_Type, Data_Rule_Type >

Powerset of a binary array.

8.51 include/barry/powerset-meat.hpp File Reference

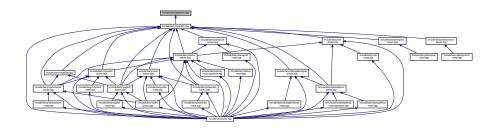
#include "powerset-bones.hpp"
Include dependency graph for powerset-meat.hpp:





8.52 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

8.52.1 Macro Definition Documentation

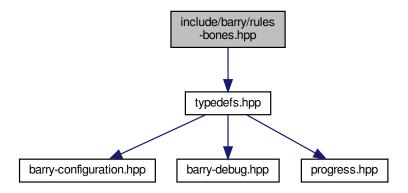
8.52.1.1 BARRY_PROGRESS_BAR_WIDTH

#define BARRY_PROGRESS_BAR_WIDTH 80

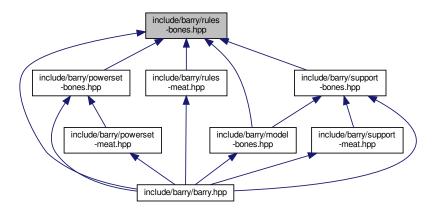
Definition at line 5 of file progress.hpp.

8.53 include/barry/rules-bones.hpp File Reference

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



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



Classes

```
    class Rule < Array_Type, Data_Type >
    Rule for determining if a cell should be included in a sequence.
```

class Rules < Array_Type, Data_Type >
 Vector of objects of class Rule.

Functions

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

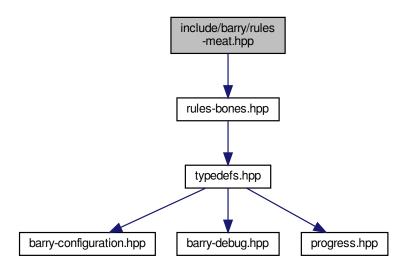
8.53.1 Function Documentation

8.53.1.1 rule_fun_default()

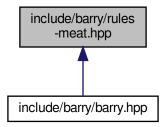
Definition at line 7 of file rules-bones.hpp.

8.54 include/barry/rules-meat.hpp File Reference

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



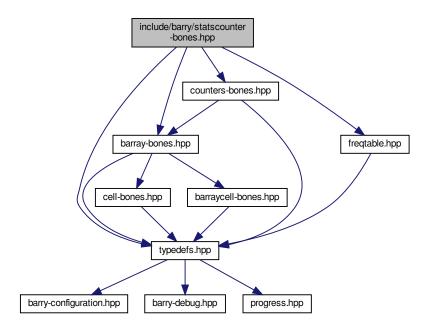
This graph shows which files directly or indirectly include this file:



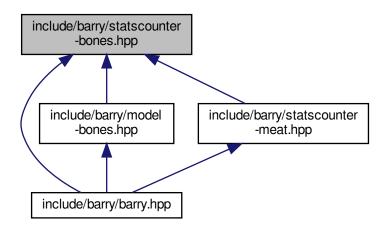
8.55 include/barry/statscounter-bones.hpp File Reference

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

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



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

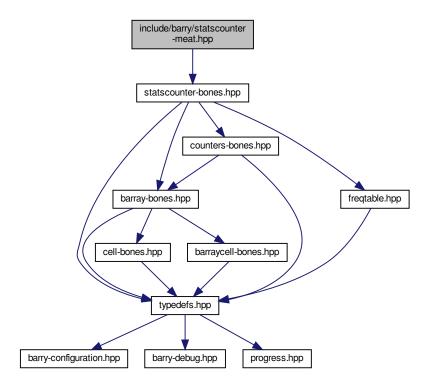


Classes

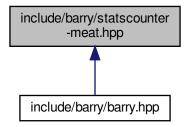
class StatsCounter < Array_Type, Data_Type >
 Count stats for a single Array.

8.56 include/barry/statscounter-meat.hpp File Reference

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



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



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)(uint i
- current_stats resize (counters->size(), 0.0)
- for (uint n=0u;n< counters->size();++n) current stats[n]
- STATSCOUNTER_TEMPLATE (void, count_current)(uint i
- STATSCOUNTER_TEMPLATE (std::vector< std::string >, get_names)() const
- STATSCOUNTER_TEMPLATE (std::vector< std::string >, get_descriptions)() const

Variables

- · Data_Type & counter
- EmptyArray = *Array
- current stats = counter.current stats
- counters = new Counters<Array_Type,Data_Type>((*counter.counters))
- counter_deleted = false
- Data_Type f_
- return
- Data Type * counters
- uint j

8.56.1 Macro Definition Documentation

8.56.1.1 STATSCOUNTER_TEMPLATE

Definition at line 11 of file statscounter-meat.hpp.

8.56.1.2 STATSCOUNTER TEMPLATE ARGS

```
template STATSCOUNTER_TEMPLATE_ARGS() <typename Array_Type</pre>, typename Data_Type>
```

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

8.56.1.3 STATSCOUNTER_TYPE

```
template Data_Type * STATSCOUNTER_TYPE( ) StatsCounter<Array_Type,Data_Type>
```

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

8.56.2 Function Documentation

8.56.2.1 clear()

```
EmptyArray clear ( )
```

8.56.2.2 for()

8.56.2.3 resize()

8.56.2.4 STATSCOUNTER_TEMPLATE() [1/9]

```
STATSCOUNTER_TEMPLATE (
StatsCounter ) const
```

8.56.2.5 STATSCOUNTER_TEMPLATE() [2/9]

```
STATSCOUNTER_TEMPLATE ( \sim StatsCounter )
```

Definition at line 30 of file statscounter-meat.hpp.

8.56.2.6 STATSCOUNTER_TEMPLATE() [3/9]

```
STATSCOUNTER_TEMPLATE (
          std::vector< std::string > ,
          get_descriptions ) const
```

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

8.56.2.7 STATSCOUNTER_TEMPLATE() [4/9]

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

8.56.2.8 STATSCOUNTER_TEMPLATE() [5/9]

8.56.2.9 STATSCOUNTER_TEMPLATE() [6/9]

8.56.2.10 STATSCOUNTER_TEMPLATE() [7/9]

8.56.2.11 STATSCOUNTER_TEMPLATE() [8/9]

Definition at line 37 of file statscounter-meat.hpp.

8.56.2.12 STATSCOUNTER_TEMPLATE() [9/9]

8.56.3 Variable Documentation

8.56.3.1 counter

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

8.56.3.2 counter_deleted

```
counter_deleted = false
```

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

8.56.3.3 counters

```
counters = new Counters<Array_Type, Data_Type>((*counter.counters))
```

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

8.56.3.4 counters_

```
Data_Type* counters_
Initial value:
{
    if (!counter_deleted)
```

Definition at line 56 of file statscounter-meat.hpp.

8.56.3.5 current_stats

```
current_stats = counter.current_stats
```

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

8.56.3.6 EmptyArray

```
EmptyArray = *Array
```

Definition at line 20 of file statscounter-meat.hpp.

```
8.56.3.7 f_
```

```
Data_Rule_Dyn_Type f_
```

Initial value:

```
counters->add_counter(f_)
```

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

8.56.3.8 j

```
uint j
```

Initial value:

```
if (counters->size() == 0u)
    throw std::logic_error("No counters added: Cannot count without knowning what to count!")
```

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

8.56.3.9 return

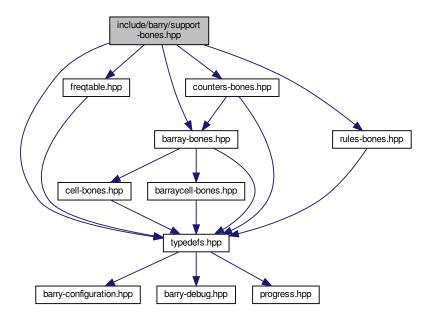
return

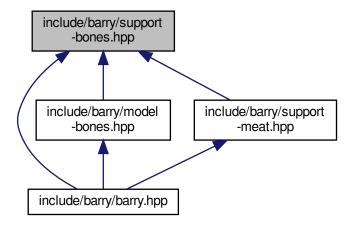
Definition at line 52 of file statscounter-meat.hpp.

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

```
#include "typedefs.hpp"
#include "barray-bones.hpp"
#include "freqtable.hpp"
#include "counters-bones.hpp"
#include "rules-bones.hpp"
```

Include dependency graph for support-bones.hpp:



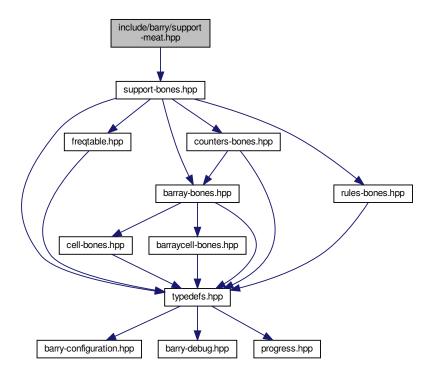


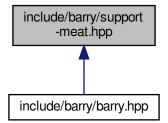
Classes

class Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >
 Compute the support of sufficient statistics.

8.58 include/barry/support-meat.hpp File Reference

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





Macros

- #define BARRY_SUPPORT_MEAT_HPP 1
- #define SUPPORT TEMPLATE ARGS()
- #define SUPPORT TYPE()
- #define SUPPORT_TEMPLATE(a, b)

Functions

- SUPPORT_TEMPLATE (void, init_support)(std
- SUPPORT TEMPLATE (void, reset array)()
- SUPPORT_TEMPLATE (void, reset_array)(const Array_Type &Array_)
- SUPPORT TEMPLATE (void, calc backend sparse)(uint pos
- calc_backend_sparse (pos+1u, array_bank, stats_bank)
- EmptyArray insert_cell (coord_i, coord_j, EmptyArray.default_val().value, false, false)
- for (uint n=0u;n< n_counters;++n)
- if (rules dyn->size() > 0u)
- if (array_bank !=nullptr) array_bank -> push_back(EmptyArray)
- EmptyArray rm cell (coord i, coord j, false, false)
- if (change stats different > 0u)
- SUPPORT TEMPLATE (void, calc backend dense)(uint pos
- calc_backend_dense (pos+1u, array_bank, stats_bank)
- EmptyArray insert cell (coord i, coord j, 1, false, false)
- SUPPORT_TEMPLATE (void, calc)(std
- SUPPORT TEMPLATE (void, add counter)(Counter< Array Type
- SUPPORT_TEMPLATE (void, set_counters)(Counters< Array_Type
- SUPPORT_TEMPLATE (void, add_rule)(Rule < Array_Type
- SUPPORT TEMPLATE (void, set rules)(Rules< Array Type
- SUPPORT_TEMPLATE (void, add_rule_dyn)(Rule < Array_Type
- SUPPORT_TEMPLATE (void, set_rules_dyn)(Rules< Array_Type
- SUPPORT_TEMPLATE (bool, eval_rules_dyn)(const std
- SUPPORT_TEMPLATE (std::vector< double >, get_counts)() const
- SUPPORT_TEMPLATE (std::vector< double > *, get_current_stats)()
- SUPPORT_TEMPLATE (void, print)() const
- SUPPORT TEMPLATE (const FreqTable <> &, get data)() const

Variables

- std::vector< Array_Type > * array_bank
- std::vector< Array Type > std::vector< double > * stats bank
- const size t & coord i = coordinates free[pos * 2u]
- const size_t & coord_j = coordinates_free[pos * 2u + 1u]
- · double tmp_chng
- unsigned int change stats different = hashes initialized[pos] ? Ou : 1u
- else
- · & hashes [pos]
- return
- Data_Counter_Type f_
- Data_Counter_Type * counters_
- delete_counters = false
- counters = counters_
- Data_Rule_Type * rules_
- delete rules = false
- rules = rules
- delete_rules_dyn = false
- rules_dyn = rules_

8.58.1 Macro Definition Documentation

8.58.1.1 BARRY_SUPPORT_MEAT_HPP

```
#define BARRY_SUPPORT_MEAT_HPP 1
```

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

8.58.1.2 SUPPORT_TEMPLATE

Value:

```
template SUPPORT_TEMPLATE_ARGS() \
inline a SUPPORT_TYPE()::b
```

Definition at line 12 of file support-meat.hpp.

8.58.1.3 SUPPORT_TEMPLATE_ARGS

```
template SUPPORT_TEMPLATE_ARGS( )
```

Value:

```
<tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre>
```

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

8.58.1.4 SUPPORT_TYPE

```
template Data_Rule_Dyn_Type * SUPPORT_TYPE( )
```

Value:

```
Support<Array_Type,Data_Counter_Type,Data_Rule_Type,\
Data_Rule_Dyn_Type>
```

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

8.58.2 Function Documentation

8.58.2.1 calc_backend_dense()

```
calc_backend_dense (
    pos+ 1u,
    array_bank ,
    stats_bank )
```

8.58.2.2 calc_backend_sparse()

```
calc_backend_sparse ( pos+\ 1u, array\_bank\ , stats\_bank\ )
```

8.58.2.3 for()

```
for ( )
```

Definition at line 162 of file support-meat.hpp.

8.58.2.4 if() [1/3]

```
if (
     array_bank ! = nullptr ) -> push_back(EmptyArray)
```

8.58.2.5 if() [2/3]

```
if ( \label{eq:change_stats_different} \mbox{,} \\ \mbox{Ou )}
```

Definition at line 242 of file support-meat.hpp.

8.58.2.6 if() [3/3]

```
if (
    rules_dyn-> size(),
    0u )
```

Definition at line 190 of file support-meat.hpp.

8.58.2.7 insert_cell() [1/2]

8.58.2.8 insert_cell() [2/2]

8.58.2.9 rm_cell()

8.58.2.10 SUPPORT_TEMPLATE() [1/17]

```
SUPPORT_TEMPLATE ( bool , eval_rules_dyn ) const
```

Definition at line 491 of file support-meat.hpp.

8.58.2.11 SUPPORT_TEMPLATE() [2/17]

Definition at line 560 of file support-meat.hpp.

8.58.2.12 SUPPORT_TEMPLATE() [3/17]

```
SUPPORT_TEMPLATE (
          std::vector< double > * ,
          get_current_stats )
```

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

8.58.2.13 SUPPORT_TEMPLATE() [4/17]

```
SUPPORT_TEMPLATE (
          std::vector< double > ,
          get_counts ) const
```

Definition at line 533 of file support-meat.hpp.

8.58.2.14 SUPPORT_TEMPLATE() [5/17]

```
SUPPORT_TEMPLATE (
     void ,
     add_counter )
```

8.58.2.15 SUPPORT_TEMPLATE() [6/17]

8.58.2.16 SUPPORT_TEMPLATE() [7/17]

8.58.2.17 SUPPORT_TEMPLATE() [8/17]

```
SUPPORT_TEMPLATE (
     void ,
     calc )
```

Definition at line 374 of file support-meat.hpp.

8.58.2.18 SUPPORT_TEMPLATE() [9/17]

8.58.2.19 SUPPORT_TEMPLATE() [10/17]

8.58.2.20 SUPPORT_TEMPLATE() [11/17]

Definition at line 16 of file support-meat.hpp.

8.58.2.21 SUPPORT_TEMPLATE() [12/17]

```
SUPPORT_TEMPLATE (
     void ,
     print ) const
```

Definition at line 549 of file support-meat.hpp.

8.58.2.22 SUPPORT_TEMPLATE() [13/17]

```
SUPPORT_TEMPLATE (
     void ,
     reset_array )
```

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

8.58.2.23 SUPPORT_TEMPLATE() [14/17]

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

8.58.2.24 SUPPORT_TEMPLATE() [15/17]

8.58.2.25 SUPPORT_TEMPLATE() [16/17]

```
SUPPORT_TEMPLATE (
     void ,
     set_rules )
```

8.58.2.26 SUPPORT_TEMPLATE() [17/17]

8.58.3 Variable Documentation

8.58.3.1 array_bank

```
std::vector< Array_Type > * array_bank
```

Definition at line 134 of file support-meat.hpp.

8.58.3.2 change_stats_different

```
unsigned int change_stats_different = hashes_initialized[pos] ? Ou : 1u
```

Definition at line 161 of file support-meat.hpp.

8.58.3.3 coord_i

```
const size_t & coord_i = coordinates_free[pos * 2u]
```

Definition at line 147 of file support-meat.hpp.

8.58.3.4 coord_j

```
const size_t & coord_j = coordinates_free[pos * 2u + 1u]
```

Definition at line 148 of file support-meat.hpp.

8.58.3.5 counters

```
counters = counters_
```

Definition at line 419 of file support-meat.hpp.

8.58.3.6 counters_

```
Data_Counter_Type* counters_
Initial value:
{
```

if (delete_counters)
 delete counters

Definition at line 412 of file support-meat.hpp.

8.58.3.7 delete_counters

```
delete_counters = false
```

Definition at line 418 of file support-meat.hpp.

8.58.3.8 delete_rules

```
delete_rules = false
```

Definition at line 452 of file support-meat.hpp.

8.58.3.9 delete_rules_dyn

```
delete_rules_dyn = false
```

Definition at line 484 of file support-meat.hpp.

8.58.3.10 else

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

8.58.3.11 f_

```
Data_Rule_Dyn_Type f_
Initial value:
{
    counters->add_counter(f_)
```

Definition at line 403 of file support-meat.hpp.

8.58.3.12 hashes

```
& hashes
```

Definition at line 220 of file support-meat.hpp.

8.58.3.13 return

```
return
```

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

8.58.3.14 rules

```
rules = rules_
```

Definition at line 453 of file support-meat.hpp.

8.58.3.15 rules_

```
Data_Rule_Dyn_Type * rules_
Initial value:
{
    if (delete_rules)
        delete rules
```

Definition at line 446 of file support-meat.hpp.

8.58.3.16 rules_dyn

```
rules_dyn = rules_
```

Definition at line 485 of file support-meat.hpp.

8.58.3.17 stats_bank

```
std::vector< Array_Type > std::vector< double > * stats_bank

Initial value:
{
    if (pos >= coordiantes_n_free)
        return
```

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

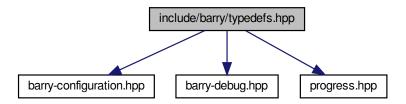
8.58.3.18 tmp_chng

double tmp_chng

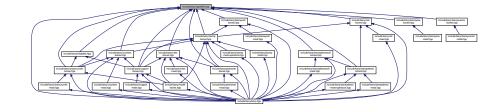
Definition at line 160 of file support-meat.hpp.

8.59 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< T >

Namespaces

• CHECK

Integer constants used to specify which cell should be check.

• EXISTS

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

Typedefs

```
typedef unsigned int uint
typedef std::vector< std::pair< std::vector< double >, uint > > Counts_type
template<typename Cell_Type >
using Row_type = Map< uint, Cell< Cell_Type >>
template<typename Cell_Type >
using Col_type = Map< uint, Cell< Cell_Type > *>
template<typename Ta = double, typename Tb = uint>
using MapVec_type = std::unordered_map< std::vector< Ta >, Tb, vecHasher< Ta >>
template<typename Array_Type , typename Data_Type >
using Counter_fun_type = std::function< double(const Array_Type &, uint, uint, Data_Type &)>
Counter and rule functions.
template<typename Array_Type , typename Data_Type >
using Rule_fun_type = std::function< bool(const Array_Type &, uint, uint, Data_Type &)>
```

Functions

```
template < typename T >
        T vec_inner_prod (const 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
```

8.59.1 Typedef Documentation

8.59.1.1 Col_type

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

Definition at line 71 of file typedefs.hpp.

8.59.1.2 Counter_fun_type

```
template<typename Array_Type , typename Data_Type >
using Counter_fun_type = std::function<double(const Array_Type &, uint, uint, Data_Type &)>
```

Counter and rule functions.

Parameters

Array_Type	a BArray
unit,uint	Focal cell
Data_Type	Data associated with the function, for example, id of the attribute in the Array.

Returns

```
Counter_fun_type a double (the change statistic)
Rule_fun_type a bool. True if the cell is blocked.
```

Definition at line 148 of file typedefs.hpp.

8.59.1.3 Counts_type

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

Definition at line 52 of file typedefs.hpp.

8.59.1.4 MapVec_type

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

Definition at line 129 of file typedefs.hpp.

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8.59.1.5 Row_type

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

Definition at line 68 of file typedefs.hpp.

8.59.1.6 Rule_fun_type

```
template<typename Array_Type , typename Data_Type >
using Rule_fun_type = std::function<bool(const Array_Type &, uint, uint, Data_Type &)>
```

Definition at line 151 of file typedefs.hpp.

8.59.1.7 uint

```
typedef unsigned int uint
```

Definition at line 18 of file typedefs.hpp.

8.59.2 Function Documentation

8.59.2.1 vec_equal()

Compares if -a- and -b- are equal.

Parameters

```
a,b Two vectors of the same length
```

Returns

 $\verb|true| if all elements are equal.$

Definition at line 162 of file typedefs.hpp.

8.59.2.2 vec_equal_approx()

Definition at line 180 of file typedefs.hpp.

8.59.2.3 vec_inner_prod() [1/2]

Definition at line 226 of file typedefs.hpp.

8.59.2.4 vec_inner_prod() [2/2]

Definition at line 203 of file typedefs.hpp.

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