barry: Your go-to motif accountant 0.0-1

Generated by Doxygen 1.9.1

1 Main Page	1
2 Module Index	3
2.1 Modules	3
3 Class Index	5
3.1 Class List	5
4 File Index	7
4.1 File List	7
5 Module Documentation	9
5.1 Counting	9
5.1.1 Detailed Description	9
5.2 Statistical Models	9
5.2.1 Detailed Description	10
5.3 Network counters	10
5.3.1 Detailed Description	10
5.3.2 Function Documentation	11
5.3.2.1 counter_absdiff()	11
5.3.2.2 counter_ctriads()	11
5.3.2.3 counter_degree()	11
5.3.2.4 counter_density()	11
5.3.2.5 counter_diff()	12
5.3.2.6 counter_edges()	12
5.3.2.7 counter_idegree()	12
5.3.2.8 counter_idegree15()	12
5.3.2.9 counter isolates()	12
5.3.2.10 counter_istar2()	13
5.3.2.11 counter_mutual()	13
5.3.2.12 counter_nodecov()	13
5.3.2.13 counter_nodeicov()	13
5.3.2.14 counter_nodematch()	13
5.3.2.15 counter_nodeocov()	14
5.3.2.16 counter_odegree()	14
5.3.2.17 counter_odegree15()	14
5.3.2.18 counter_ostar2()	14
5.3.2.19 counter_ttriads()	14
5.3.2.20 NETWORK_COUNTER()	15
5.4 Phylo counters	15
5.4.1 Detailed Description	15
5.4.2 Function Documentation	16
5.4.2 Function Documentation	16
5.4.2.2 counter_cogain()	16

	5.4.2.3 counter_gains()	1/
	5.4.2.4 counter_gains_k_offspring()	17
	5.4.2.5 counter_genes_changing()	17
	5.4.2.6 counter_longest()	17
	5.4.2.7 counter_loss()	18
	5.4.2.8 counter_maxfuns()	18
	5.4.2.9 counter_neofun()	18
	5.4.2.10 counter_neofun_a2b()	18
	5.4.2.11 counter_overall_changes()	19
	5.4.2.12 counter_overall_gains()	19
	5.4.2.13 counter_overall_loss()	19
	5.4.2.14 counter_subfun()	19
	5.5 Phylo rules	20
	5.5.1 Detailed Description	20
	5.5.2 Function Documentation	20
	5.5.2.1 rule_dyn_limit_changes()	20
6	Namespace Documentation	23
	6.1 barry Namespace Reference	23
	6.1.1 Detailed Description	23
	6.2 barry::counters Namespace Reference	23
	6.2.1 Detailed Description	23
	6.3 barry::counters::network Namespace Reference	24
	6.4 barry::counters::phylo Namespace Reference	24
	6.5 CHECK Namespace Reference	24
	6.5.1 Detailed Description	24
	6.5.2 Variable Documentation	24
	6.5.2.1 BOTH	24
	6.5.2.2 NONE	24
	6.5.2.3 ONE	24
	6.5.2.4 TWO	25
	6.6 EXISTS Namespace Reference	25
	6.6.1 Detailed Description	25
	6.6.2 Variable Documentation	25
	6.6.2.1 AS_ONE	25
	6.6.2.2 AS_ZERO	25
	6.6.2.3 BOTH	26
	6.6.2.4 NONE	26
	6.6.2.5 ONE	26
	6.6.2.6 TWO	26
	6.6.2.7 UKNOWN	26
7	Class Desumentation	07
1	Class Documentation	27

7.1 BArray< Cell_Type, Data_Type > Class Template Reference	27
7.1.1 Detailed Description	29
7.1.2 Constructor & Destructor Documentation	30
7.1.2.1 BArray() [1/6]	30
7.1.2.2 BArray() [2/6]	30
<b>7.1.2.3 BArray()</b> [3/6]	30
7.1.2.4 BArray() [4/6]	31
<b>7.1.2.5 BArray()</b> [5/6]	31
7.1.2.6 BArray() [6/6]	31
7.1.2.7 ∼BArray()	31
7.1.3 Member Function Documentation	31
7.1.3.1 clear()	31
7.1.3.2 col()	32
7.1.3.3 D() [1/2]	32
7.1.3.4 D() [2/2]	32
7.1.3.5 default_val()	32
7.1.3.6 get_cell()	32
7.1.3.7 get_col()	32
7.1.3.8 get_col_vec() [1/2]	33
7.1.3.9 get_col_vec() [2/2]	33
7.1.3.10 get_entries()	33
7.1.3.11 get_row()	33
7.1.3.12 get_row_vec() [1/2]	33
7.1.3.13 get_row_vec() [2/2]	34
7.1.3.14 insert_cell() [1/3]	34
7.1.3.15 insert_cell() [2/3]	34
7.1.3.16 insert_cell() [3/3]	34
7.1.3.17 is_empty()	34
7.1.3.18 ncol()	35
7.1.3.19 nnozero()	35
7.1.3.20 nrow()	35
7.1.3.21 operator()() [1/2]	35
7.1.3.22 operator()() [2/2]	35
7.1.3.23 operator*=()	35
7.1.3.24 operator+=() [1/3]	36
7.1.3.25 operator+=() [2/3]	36
7.1.3.26 operator+=() [3/3]	36
<b>7.1.3.27 operator-=()</b> [1/3]	36
<b>7.1.3.28 operator-=()</b> [2/3]	36
<b>7.1.3.29 operator-=()</b> [3/3]	36
7.1.3.30 operator/=()	37
7.1.3.31 operator=() [1/2]	37

7.1.3.32 operator=() [2/2]	 . 37
7.1.3.33 operator==()	 . 37
7.1.3.34 out_of_range()	 . 37
7.1.3.35 print()	 . 37
7.1.3.36 reserve()	 . 38
7.1.3.37 resize()	 . 38
7.1.3.38 rm_cell()	 . 38
7.1.3.39 row()	 . 38
7.1.3.40 set_data()	 . 38
7.1.3.41 swap_cells()	 . 39
7.1.3.42 swap_cols()	 . 39
7.1.3.43 swap_rows()	 . 39
7.1.3.44 toggle_cell()	 . 39
7.1.3.45 toggle_lock()	 . 40
7.1.3.46 transpose()	 . 40
7.1.3.47 zero_col()	 . 40
7.1.3.48 zero_row()	 . 40
7.1.4 Friends And Related Function Documentation	 . 40
7.1.4.1 BArrayCell < Cell_Type, Data_Type >	 . 40
7.1.4.2 BArrayCell_const< Cell_Type, Data_Type >	 . 41
7.1.5 Member Data Documentation	 . 41
7.1.5.1 visited	 . 41
7.2 BArrayCell< Cell_Type, Data_Type > Class Template Reference	 . 41
7.2.1 Detailed Description	 . 41
7.2.2 Constructor & Destructor Documentation	 . 42
7.2.2.1 BArrayCell()	 . 42
7.2.2.2 ~BArrayCell()	 . 42
7.2.3 Member Function Documentation	 . 42
7.2.3.1 operator Cell_Type()	 . 42
7.2.3.2 operator*=()	 . 42
7.2.3.3 operator+=()	 . 43
7.2.3.4 operator-=()	 . 43
7.2.3.5 operator/=()	 . 43
7.2.3.6 operator=()	 . 43
7.2.3.7 operator==()	 . 43
7.3 BArrayCell_const< Cell_Type, Data_Type > Class Template Reference	 . 44
7.3.1 Detailed Description	 . 44
7.3.2 Constructor & Destructor Documentation	 . 44
7.3.2.1 BArrayCell_const()	 . 44
7.3.2.2 ~BArrayCell_const()	 . 44
7.3.3 Member Function Documentation	 . 45
7.3.3.1 operator Cell_Type()	 . 45

7.3.3.2 operator"!=()	45
7.3.3.3 operator<()	45
7.3.3.4 operator<=()	45
7.3.3.5 operator==()	45
7.3.3.6 operator>()	46
7.3.3.7 operator>=()	46
7.4 BArrayCol< Cell_Type, Data_Type > Class Template Reference	46
7.4.1 Detailed Description	46
7.4.2 Constructor & Destructor Documentation	47
7.4.2.1 BArrayCol()	47
7.4.2.2 ~BArrayCol()	47
7.4.3 Member Function Documentation	47
7.4.3.1 begin()	47
7.4.3.2 end()	47
7.4.3.3 operator Cell_Type()	48
7.4.3.4 operator*=()	48
7.4.3.5 operator+=()	48
7.4.3.6 operator-=()	48
7.4.3.7 operator/=()	48
7.4.3.8 operator=()	49
7.4.3.9 operator==()	49
7.5 BArrayCol_const< Cell_Type, Data_Type > Class Template Reference	49
7.5.1 Detailed Description	49
7.5.2 Constructor & Destructor Documentation	50
7.5.2.1 ~BArrayCol_const()	50
7.5.3 Member Function Documentation	50
7.5.3.1 BArrayCol()	50
7.5.3.2 operator"!=()	50
7.5.3.3 operator<()	50
7.5.3.4 operator<=()	51
7.5.3.5 operator==()	51
7.5.3.6 operator>()	51
7.5.3.7 operator>=()	51
7.6 BArrayDense< Cell_Type, Data_Type > Class Template Reference	51
7.6.1 Detailed Description	52
7.6.2 Constructor & Destructor Documentation	52
7.6.2.1 BArrayDense() [1/2]	52
7.6.2.2 BArrayDense() [2/2]	52
7.6.2.3 ~BArrayDense()	53
7.6.3 Member Function Documentation	53
7.6.3.1 elements_ptr()	53
7.6.3.2 elements_raw()	53

7.6.3.3 fill()	 . 53
7.6.3.4 ncol()	 . 53
7.6.3.5 nrow()	 . 54
7.6.3.6 operator()()	 . 54
7.6.3.7 operator[]()	 . 54
7.6.3.8 print()	 . 54
7.7 Cell < Cell_Type > Class Template Reference	 . 54
7.7.1 Detailed Description	 . 55
7.7.2 Constructor & Destructor Documentation	 . 55
<b>7.7.2.1 Cell()</b> [1/7]	 . 55
<b>7.7.2.2 Cell()</b> [2/7]	 . 56
7.7.2.3 ~Cell()	 . 56
<b>7.7.2.4 Cell()</b> [3/7]	 . 56
7.7.2.5 Cell() [4/7]	 . 56
<b>7.7.2.6 Cell()</b> [5/7]	 . 56
<b>7.7.2.7 Cell()</b> [6/7]	 . 57
<b>7.7.2.8 Cell()</b> [7/7]	 . 57
7.7.3 Member Function Documentation	 . 57
7.7.3.1 add() [1/4]	 . 57
7.7.3.2 add() [2/4]	 . 57
<b>7.7.3.3 add()</b> [3/4]	 . 57
7.7.3.4 add() [4/4]	 . 58
7.7.3.5 operator Cell_Type()	 . 58
7.7.3.6 operator=() [1/2]	 . 58
7.7.3.7 operator=() [2/2]	 . 58
7.7.4 Member Data Documentation	 . 58
7.7.4.1 value	 . 58
7.7.4.2 visited	 . 59
$7.8 \ ConstBArrayRowlter < Cell\_Type, \ Data\_Type > Class \ Template \ Reference \ . \ . \ . \ . \ . \ . \ . \ . \ . \ $	 . 59
7.8.1 Detailed Description	 . 60
7.8.2 Constructor & Destructor Documentation	 . 60
7.8.2.1 ConstBArrayRowlter()	 . 60
$7.8.2.2 \sim ConstBArrayRowlter()$	 . 60
7.8.3 Member Data Documentation	 . 60
7.8.3.1 Array	 . 60
7.8.3.2 current_col	 . 60
7.8.3.3 current_row	 . 61
7.8.3.4 iter	 . 61
7.9 Counter< Array_Type, Data_Type > Class Template Reference	 . 61
7.9.1 Detailed Description	 . 62
7.9.2 Constructor & Destructor Documentation	 . 62
7.9.2.1 Counter() [1/4]	 . 62

<b>7.9.2.2 Counter()</b> [2/4]	. 63
7.9.2.3 Counter() [3/4]	. 63
7.9.2.4 Counter() [4/4]	. 63
7.9.2.5 ~Counter()	. 63
7.9.3 Member Function Documentation	. 63
7.9.3.1 count()	. 64
7.9.3.2 init()	. 64
7.9.3.3 operator=() [1/2]	. 64
7.9.3.4 operator=() [2/2]	. 64
7.9.4 Member Data Documentation	. 64
7.9.4.1 count_fun	. 65
7.9.4.2 data	. 65
7.9.4.3 delete_data	. 65
7.9.4.4 desc	. 65
7.9.4.5 init_fun	. 65
7.9.4.6 name	. 66
7.10 Counters< Array_Type, Data_Type > Class Template Reference	. 66
7.10.1 Detailed Description	. 66
7.10.2 Constructor & Destructor Documentation	. 67
7.10.2.1 Counters() [1/3]	. 67
7.10.2.2 ~Counters()	. 67
<b>7.10.2.3 Counters()</b> [2/3]	. 67
<b>7.10.2.4 Counters()</b> [3/3]	. 67
7.10.3 Member Function Documentation	. 68
<b>7.10.3.1 add_counter()</b> [1/3]	. 68
<b>7.10.3.2 add_counter()</b> [2/3]	. 68
<b>7.10.3.3 add_counter()</b> [3/3]	. 68
7.10.3.4 clear()	. 69
7.10.3.5 operator=() [1/2]	. 69
7.10.3.6 operator=() [2/2]	. 69
7.10.3.7 operator[]()	. 70
7.10.3.8 size()	. 70
7.11 Entries < Cell_Type > Class Template Reference	. 70
7.11.1 Detailed Description	. 71
7.11.2 Constructor & Destructor Documentation	. 71
7.11.2.1 Entries() [1/2]	. 71
7.11.2.2 Entries() [2/2]	. 71
7.11.2.3 ~Entries()	. 71
7.11.3 Member Function Documentation	. 72
7.11.3.1 resize()	. 72
7.11.4 Member Data Documentation	. 72
7.11.4.1 source	72

7.11.4.2 target	72
7.11.4.3 val	72
7.12 Flock Class Reference	73
7.12.1 Detailed Description	73
7.12.2 Constructor & Destructor Documentation	74
7.12.2.1 Flock()	74
7.12.2.2 ~Flock()	74
7.12.3 Member Function Documentation	74
7.12.3.1 add_data()	74
7.12.3.2 get_counters()	75
7.12.3.3 get_support()	75
7.12.3.4 init()	75
7.12.3.5 likelihood_joint()	75
7.12.3.6 nfuns()	76
7.12.3.7 nleafs()	76
7.12.3.8 nnodes()	76
7.12.3.9 nterms()	76
7.12.3.10 ntrees()	76
7.12.3.11 operator()()	76
7.12.3.12 set_seed()	77
7.12.3.13 support_size()	77
7.12.4 Member Data Documentation	77
7.12.4.1 dat	77
7.12.4.2 initialized	78
7.12.4.3 nfunctions	78
7.12.4.4 rengine	78
7.12.4.5 support	78
7.13 FreqTable $<$ T $>$ Class Template Reference	78
7.13.1 Detailed Description	79
7.13.2 Constructor & Destructor Documentation	79
7.13.2.1 FreqTable()	79
7.13.2.2 ∼FreqTable()	79
7.13.3 Member Function Documentation	79
7.13.3.1 add()	79
7.13.3.2 as_vector()	80
7.13.3.3 clear()	80
7.13.3.4 get_data()	80
7.13.3.5 get_data_ptr()	80
7.13.3.6 print()	80
7.13.3.7 reserve()	81
7.13.3.8 size()	81
7.14 Geese Class Reference	81

7.14.1 Detailed Description
7.14.2 Constructor & Destructor Documentation
7.14.2.1 Geese() [1/4]
7.14.2.2 Geese() [2/4]
7.14.2.3 Geese() [3/4]
7.14.2.4 Geese() [4/4]
7.14.2.5 ~Geese()
7.14.3 Member Function Documentation
7.14.3.1 calc_reduced_sequence()
7.14.3.2 calc_sequence()
7.14.3.3 get_counters()
7.14.3.4 get_probabilities()
7.14.3.5 get_rengine()
7.14.3.6 get_states()
7.14.3.7 get_support()
7.14.3.8 inherit_support()
7.14.3.9 init()
7.14.3.10 init_node()
7.14.3.11 likelihood()
7.14.3.12 likelihood_exhaust()
7.14.3.13 nfuns()
7.14.3.14 nleafs()
7.14.3.15 nnodes()
7.14.3.16 nterms()
7.14.3.17 observed_counts()
7.14.3.18 operator=() [1/2]
7.14.3.19 operator=() [2/2]
7.14.3.20 predict()
7.14.3.21 predict_backend()
7.14.3.22 print_observed_counts()
7.14.3.23 set_seed()
7.14.3.24 simulate()
7.14.3.25 support_size()
7.14.3.26 update_annotations()
7.14.4 Member Data Documentation
7.14.4.1 delete_rengine
7.14.4.2 delete_support
7.14.4.3 initialized
7.14.4.4 map_to_nodes
7.14.4.5 nfunctions
7.14.4.6 nodes
7.14.4.7 reduced sequence 90

	7.14.4.8 sequence	 	 	 	 	 
	el< Array_Type, Data_Counter_Type, Data_F e Reference			 		
7.15	5.1 Detailed Description	 	 	 	 	 
7.15	5.2 Constructor & Destructor Documentation .	 	 	 	 	 
	<b>7.15.2.1 Model()</b> [1/3]	 	 	 	 	 
	7.15.2.2 Model() [2/3]	 	 	 	 	 
	<b>7.15.2.3 Model()</b> [3/3]	 	 	 	 	 
	7.15.2.4 ~Model()	 	 	 	 	 
7.15	5.3 Member Function Documentation	 	 	 	 	 
	7.15.3.1 add_array()	 	 	 	 	 
	7.15.3.2 add_counter() [1/3]	 	 	 	 	 
	7.15.3.3 add_counter() [2/3]	 	 	 	 	 
	7.15.3.4 add_counter() [3/3]	 	 	 	 	 
	<b>7.15.3.5 add_rule()</b> [1/3]	 	 	 	 	 
	<b>7.15.3.6 add_rule()</b> [2/3]	 	 	 	 	 
	<b>7.15.3.7 add_rule()</b> [3/3]	 	 	 	 	 
	7.15.3.8 add_rule_dyn() [1/3]	 	 	 	 	 
	7.15.3.9 add_rule_dyn() [2/3]	 	 	 	 	 
	<b>7.15.3.10 add_rule_dyn()</b> [3/3]	 	 	 	 	 
	7.15.3.11 get_counters()	 	 	 	 	 
	7.15.3.12 get_norm_const()	 	 	 	 	 
	7.15.3.13 get_pset()	 	 	 	 	 
	7.15.3.14 get_rengine()	 	 	 	 	 
	7.15.3.15 get_rules()	 	 	 	 	 
	7.15.3.16 get_rules_dyn()	 	 	 	 	 
	7.15.3.17 get_stats()	 	 	 	 	 
	7.15.3.18 get_support()	 	 	 	 	 
	<b>7.15.3.19 likelihood()</b> [1/3]	 	 	 	 	 
	<b>7.15.3.20 likelihood()</b> [2/3]	 	 	 	 	 
	<b>7.15.3.21 likelihood()</b> [3/3]	 	 	 	 	 
	7.15.3.22 likelihood_total()	 	 	 	 	 
	7.15.3.23 nterms()	 	 	 	 	 
	7.15.3.24 operator=()	 	 	 	 	 
	7.15.3.25 print_stats()	 	 	 	 	 
	7.15.3.26 sample() [1/2]	 	 	 	 	 
	<b>7.15.3.27 sample()</b> [2/2]	 	 	 	 	 
	7.15.3.28 set_counters()	 	 	 	 	 
	7.15.3.29 set_keygen()	 	 	 	 	 
	7.15.3.30 set_rengine()	 	 	 	 	 
	7.15.3.31 set_rules()	 	 	 	 	 
	7.15.3.32 set_rules_dyn()	 	 	 	 	 

7.15.3.33 set_seed()	101
7.15.3.34 size()	101
7.15.3.35 size_unique()	102
7.15.3.36 store_psets()	102
7.15.3.37 support_size()	102
7.16 NetCounterData Class Reference	102
7.16.1 Detailed Description	103
7.16.2 Constructor & Destructor Documentation	103
7.16.2.1 NetCounterData() [1/2]	103
7.16.2.2 NetCounterData() [2/2]	103
7.16.2.3 ∼NetCounterData()	103
7.16.3 Member Data Documentation	103
7.16.3.1 indices	104
7.16.3.2 numbers	104
7.17 NetworkData Class Reference	104
7.17.1 Detailed Description	104
7.17.2 Constructor & Destructor Documentation	105
7.17.2.1 NetworkData() [1/3]	105
7.17.2.2 NetworkData() [2/3]	105
<b>7.17.2.3 NetworkData()</b> [3/3]	105
7.17.2.4 ~NetworkData()	106
7.17.3 Member Data Documentation	106
7.17.3.1 directed	106
7.17.3.2 vertex_attr	106
7.18 Node Class Reference	106
7.18.1 Detailed Description	107
7.18.2 Constructor & Destructor Documentation	107
7.18.2.1 Node() [1/5]	108
<b>7.18.2.2 Node()</b> [2/5]	108
<b>7.18.2.3 Node()</b> [3/5]	108
<b>7.18.2.4 Node()</b> [4/5]	108
<b>7.18.2.5 Node()</b> [5/5]	108
7.18.2.6 ~Node()	109
7.18.3 Member Function Documentation	109
7.18.3.1 get_parent()	109
7.18.3.2 is_leaf()	109
7.18.4 Member Data Documentation	109
7.18.4.1 annotations	109
7.18.4.2 array	109
7.18.4.3 arrays	110
7.18.4.4 duplication	110
7.18.4.5 id	110

7.18.4.6 narray	. 110
7.18.4.7 offspring	. 110
7.18.4.8 ord	. 111
7.18.4.9 parent	. 111
7.18.4.10 probability	. 111
7.18.4.11 subtree_prob	. 111
7.18.4.12 visited	. 111
7.19 NodeData Class Reference	. 112
7.19.1 Detailed Description	. 112
7.19.2 Constructor & Destructor Documentation	. 112
7.19.2.1 NodeData() [1/2]	. 112
7.19.2.2 NodeData() [2/2]	. 112
7.19.2.3 ~NodeData()	. 113
7.19.3 Member Data Documentation	. 113
7.19.3.1 blengths	. 113
7.19.3.2 duplication	. 113
7.19.3.3 states	. 113
7.20 PhyloRuleDynData Class Reference	. 113
7.20.1 Detailed Description	. 114
7.20.2 Constructor & Destructor Documentation	. 114
7.20.2.1 PhyloRuleDynData()	. 114
7.20.2.2 ∼PhyloRuleDynData()	. 114
7.20.3 Member Data Documentation	. 114
7.20.3.1 counts	. 115
7.20.3.2 duplication	. 115
7.20.3.3 lb	. 115
7.20.3.4 pos	. 115
7.20.3.5 ub	. 115
7.21 PowerSet< Array_Type, Data_Rule_Type > Class Template Reference	. 116
7.21.1 Detailed Description	. 117
7.21.2 Constructor & Destructor Documentation	. 117
7.21.2.1 PowerSet() [1/3]	. 117
<b>7.21.2.2 PowerSet()</b> [2/3]	. 117
<b>7.21.2.3 PowerSet()</b> [3/3]	. 118
7.21.2.4 ~PowerSet()	. 118
7.21.3 Member Function Documentation	. 118
7.21.3.1 add_rule() [1/3]	. 118
7.21.3.2 add_rule() [2/3]	. 118
<b>7.21.3.3 add_rule()</b> [3/3]	. 118
7.21.3.4 begin()	. 119
7.21.3.5 calc()	. 119
7.21.3.6 end()	. 119

7.21.3.7 get_data()	119
7.21.3.8 get_data_ptr()	119
7.21.3.9 init_support()	120
7.21.3.10 operator[]()	120
7.21.3.11 reset()	120
7.21.3.12 size()	120
7.21.4 Member Data Documentation	120
7.21.4.1 coordinates_free	120
7.21.4.2 coordinates_locked	121
7.21.4.3 data	121
7.21.4.4 EmptyArray	121
7.21.4.5 M	121
7.21.4.6 N	121
7.21.4.7 rules	122
7.21.4.8 rules_deleted	122
7.22 Rule < Array_Type, Data_Type > Class Template Reference	122
7.22.1 Detailed Description	123
7.22.2 Constructor & Destructor Documentation	123
7.22.2.1 Rule() [1/2]	123
<b>7.22.2.2 Rule()</b> [2/2]	123
7.22.2.3 ~Rule()	123
7.22.3 Member Function Documentation	124
7.22.3.1 D()	124
7.22.3.2 operator()()	124
7.23 Rules< Array_Type, Data_Type > Class Template Reference	124
7.23.1 Detailed Description	125
7.23.2 Constructor & Destructor Documentation	125
7.23.2.1 Rules() [1/2]	125
7.23.2.2 Rules() [2/2]	125
7.23.2.3 ∼Rules()	126
7.23.3 Member Function Documentation	126
7.23.3.1 add_rule() [1/3]	126
7.23.3.2 add_rule() [2/3]	126
<b>7.23.3.3 add_rule()</b> [3/3]	126
7.23.3.4 clear()	126
7.23.3.5 get_seq()	126
7.23.3.6 operator()()	127
7.23.3.7 operator=()	127
7.23.3.8 size()	128
7.24 StatsCounter< Array_Type, Data_Type > Class Template Reference	128
7.24.1 Detailed Description	128
7.24.2 Constructor & Destructor Documentation	129

7.24.2.1 StatsCounter() [1/2]	129
7.24.2.2 StatsCounter() [2/2]	129
7.24.2.3 ~StatsCounter()	129
7.24.3 Member Function Documentation	129
7.24.3.1 add_counter() [1/2]	129
7.24.3.2 add_counter() [2/2]	130
7.24.3.3 count_all()	130
7.24.3.4 count_current()	130
7.24.3.5 count_init()	130
7.24.3.6 get_counters()	130
7.24.3.7 reset_array()	130
7.24.3.8 set_counters()	131
7.25 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class Tem-	
plate Reference	
7.25.1 Detailed Description	
7.25.2 Constructor & Destructor Documentation	
<b>7.25.2.1 Support()</b> [1/3]	
<b>7.25.2.2 Support()</b> [2/3]	
<b>7.25.2.3 Support()</b> [3/3]	
7.25.2.4 ~Support()	
7.25.3 Member Function Documentation	
7.25.3.1 add_counter() [1/2]	
7.25.3.2 add_counter() [2/2]	
<b>7.25.3.3 add_rule()</b> [1/2]	
<b>7.25.3.4 add_rule()</b> [2/2]	
7.25.3.5 add_rule_dyn() [1/2]	
7.25.3.6 add_rule_dyn() [2/2]	
7.25.3.7 calc()	136
7.25.3.8 get_counters()	137
7.25.3.9 get_counts()	137
7.25.3.10 get_counts_ptr()	137
7.25.3.11 get_current_stats()	137
7.25.3.12 get_data()	138
7.25.3.13 get_rules()	138
7.25.3.14 get_rules_dyn()	138
7.25.3.15 init_support()	138
7.25.3.16 print()	139
7.25.3.17 reset_array() [1/2]	139
7.25.3.18 reset_array() [2/2]	139
7.25.3.19 set_counters()	139
7.25.3.20 set_rules()	139
7.25.3.21 set_rules_dyn()	140

 . 140
 . 140
 . 140
 . 140
 . 141
 . 142
143
_
. 153

8.10.1.2 BARRY_ISFINITE
8.10.1.3 BARRY_MAX_NUM_ELEMENTS
8.10.1.4 BARRY_SAFE_EXP
8.10.1.5 printf_barry
8.10.2 Typedef Documentation
8.10.2.1 Map
8.11 include/barry/barry.hpp File Reference
8.11.1 Macro Definition Documentation
8.11.1.1 BARRY_HPP
8.11.1.2 BARRY_VERSION
8.11.1.3 COUNTER_FUNCTION
8.11.1.4 COUNTER_LAMBDA
8.11.1.5 RULE_FUNCTION
8.11.1.6 RULE_LAMBDA
8.12 include/barry/cell-bones.hpp File Reference
8.13 include/barry/cell-meat.hpp File Reference
8.14 include/barry/col-bones.hpp File Reference
8.15 include/barry/counters-bones.hpp File Reference
8.16 include/barry/counters-meat.hpp File Reference
8.17 include/barry/counters/network.hpp File Reference
8.17.1 Macro Definition Documentation
8.17.1.1 NET_C_DATA_IDX
8.17.1.2 NET_C_DATA_NUM
8.17.1.3 NETWORK_COUNTER
8.17.1.4 NETWORK_COUNTER_LAMBDA
8.17.1.5 NETWORK_RULE
8.17.1.6 NETWORK_RULE_LAMBDA
8.17.2 Typedef Documentation
8.17.2.1 NetCounter
8.17.2.2 NetCounters
8.17.2.3 NetModel
8.17.2.4 NetRule
8.17.2.5 NetRules
8.17.2.6 NetStatsCounter
8.17.2.7 NetSupport
8.17.2.8 Network
8.17.3 Function Documentation
8.17.3.1 rules_zerodiag()
8.18 include/barry/counters/phylo.hpp File Reference
8.18.1 Macro Definition Documentation
8.18.1.1 PHYLO_CHECK_MISSING
8.18.1.2 PHYLO_COUNTER_LAMBDA

8.18.1.3 PHYLO_RULE_DYN_LAMBDA
8.18.2 Typedef Documentation
8.18.2.1 PhyloArray
8.18.2.2 PhyloCounter
8.18.2.3 PhyloCounterData
8.18.2.4 PhyloCounters
8.18.2.5 PhyloModel
8.18.2.6 PhyloPowerSet
8.18.2.7 PhyloRule
8.18.2.8 PhyloRuleData
8.18.2.9 PhyloRuleDyn
8.18.2.10 PhyloRules
8.18.2.11 PhyloRulesDyn
8.18.2.12 PhyloStatsCounter
8.18.2.13 PhyloSupport
8.18.3 Function Documentation
8.18.3.1 get_last_name()
8.19 include/barry/model-bones.hpp File Reference
8.19.1 Function Documentation
8.19.1.1 keygen_default()
8.19.1.2 likelihood_()
8.19.1.3 update_normalizing_constant()
8.20 include/barry/model-meat.hpp File Reference
8.21 include/barry/models/geese.hpp File Reference
8.22 include/barry/models/geese/flock-bones.hpp File Reference
8.23 include/barry/models/geese/flock-meet.hpp File Reference
8.24 include/barry/models/geese/geese-bones.hpp File Reference
8.24.1 Macro Definition Documentation
8.24.1.1 INITIALIZED
8.24.2 Function Documentation
8.24.2.1 keygen_full()
8.24.2.2 RULE_FUNCTION()
8.24.2.3 vec_diff()
8.24.2.4 vector_caster()
8.25 include/barry/models/geese/geese-meat-constructors.hpp File Reference
8.26 include/barry/models/geese/geese-meat-likelihood.hpp File Reference
8.27 include/barry/models/geese/geese-meat-likelihood_exhaust.hpp File Reference
8.28 include/barry/models/geese/geese-meat-predict.hpp File Reference
8.29 include/barry/models/geese/geese-meat-simulate.hpp File Reference
8.30 include/barry/models/geese/geese-meat.hpp File Reference
8.31 include/barry/models/geese/geese-node-bones.hpp File Reference
8.32 include/barry/powerset-bones hop File Reference

Index

8.33 include/barry/powerset-meat.hpp File Reference
8.34 include/barry/rules-bones.hpp File Reference
8.34.1 Function Documentation
8.34.1.1 rule_fun_default()
8.35 include/barry/rules-meat.hpp File Reference
8.36 include/barry/statscounter-bones.hpp File Reference
8.37 include/barry/statscounter-meat.hpp File Reference
8.38 include/barry/statsdb.hpp File Reference
8.39 include/barry/support-bones.hpp File Reference
8.40 include/barry/support-meat.hpp File Reference
8.40.1 Macro Definition Documentation
8.40.1.1 BARRY_SUPPORT_MEAT_HPP
8.41 include/barry/typedefs.hpp File Reference
8.41.1 Typedef Documentation
8.41.1.1 Col_type
8.41.1.2 Counter_fun_type
8.41.1.3 Counts_type
8.41.1.4 MapVec_type
8.41.1.5 Row_type
8.41.1.6 Rule_fun_type
8.41.1.7 uint
8.41.2 Function Documentation
8.41.2.1 vec_equal()
8.41.2.2 vec_equal_approx()
8.41.2.3 vec_inner_prod()
8.42 README.md File Reference

199

## **Main Page**

### Barry: your to-go motif accountant

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

### **Examples**

#### Counting statistics in a graph

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

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

2 Main Page

```
netcounters::counter_ttriads(counter.counters);
  netcounters::counter_isolates(counter.counters);
  netcounters::counter_ctriads(counter.counters);
  netcounters::counter_mutual(counter.counters);
  // Counting and printing the results
  std::vector< double > counts = counter.count_all();
  std::cout «
    "Edges : " « counts[0] « std::endl «
"Transitive triads : " « counts[1] « std::endl «
"Isolates : " « counts[2] « std::endl «
"C triads : " « counts[3] « std::endl «
"Mutuals : " « counts[4] « std::endl;
 return 0;
Compiling this program using g++
g++ -std=c++11 -Wall -pedantic 08-counts.cpp -o counts && ./counts
Yields the following output:
Current view
                1
  0,] 1 1
   1,] .
            1
  . 1
   5,]
New view
  0,] .
1,] 1
            1 1
  1,] 1 .
2,] 1 .
                    . 1
   3,] .
4,] 1
           . 1 . .
   5,] . . . .
Edges
Transitive triads : 3
Isolates
C triads
```

#### **Code of Conduct**

Mutuals

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

# **Module Index**

### 2.1 Modules

Here is a list of all modules:

Counting																							9
Statistical Models																							ç
Network counters																							10
Phylo counters																							15
Phylo rules																							20

4 Module Index

# **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	27
BArrayCell< Cell_Type, Data_Type >	41
BArrayCell_const< Cell_Type, Data_Type >	44
BArrayCol < Cell_Type, Data_Type >	46
BArrayCol_const< Cell_Type, Data_Type >	49
BArrayDense< Cell_Type, Data_Type >	
Dense bi-dimensional array	51
Cell< Cell_Type >	
Entries in BArray. For now, it only has two members:	54
ConstBArrayRowlter < Cell_Type, Data_Type >	59
Counter< Array_Type, Data_Type >	
A counter function based on change statistics	61
Counters< Array_Type, Data_Type >	
Vector of counters	66
Entries < Cell_Type >	
A wrapper class to store source, target, val from a BArray object	70
Flock	
A Flock is a group of Geese	73
FreqTable < T >	
Database of statistics	78
Geese	
Annotated Phylo Model	81
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:	90
NetCounterData	
Data class used to store arbitrary uint or double vectors	102
NetworkData	
Data class for Networks	104
Node	
A single node for the model	106
NodeData	
Data definition for the PhyloArray class	112
PhyloRuleDynData	113

6 Class Index

PowerSet < Array_Type, Data_Rule_Type >
Powerset of a binary array
Rule < Array_Type, Data_Type >
Rule for determining if a cell should be included in a sequence
Rules < Array_Type, Data_Type >
Vector of objects of class Rule
StatsCounter< Array_Type, Data_Type >
Count stats for a single Array
Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >
Compute the support of sufficient statistics
vecHasher< T >

# 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/barraycol-bones.hpp
include/barry/barraycol-meat.hpp
include/barry/barraydense-bones.hpp
include/barry/barry-configuration.hpp
include/barry/barry.hpp
include/barry/cell-bones.hpp
include/barry/cell-meat.hpp
include/barry/col-bones.hpp
include/barry/counters-bones.hpp
include/barry/counters-meat.hpp
include/barry/model-bones.hpp
include/barry/model-meat.hpp
include/barry/powerset-bones.hpp
include/barry/powerset-meat.hpp
include/barry/rules-bones.hpp
include/barry/rules-meat.hpp
include/barry/statscounter-bones.hpp
include/barry/statscounter-meat.hpp
include/barry/statsdb.hpp
include/barry/support-bones.hpp
include/barry/support-meat.hpp
include/barry/typedefs.hpp
include/barry/counters/network.hpp
include/barry/counters/phylo.hpp
include/barry/models/geese.hpp
include/barry/models/geese/flock-bones.hpp
include/barry/models/geese/flock-meet.hpp
include/barry/models/geese/geese-bones.hpp
include/barry/models/geese/geese-meat-constructors.hpp

8 File Index

include/barry/models/geese/geese-meat-likelihood.hpp	179
include/barry/models/geese/geese-meat-likelihood_exhaust.hpp	180
include/barry/models/geese/geese-meat-predict.hpp	181
include/barry/models/geese/geese-meat-simulate.hpp	181
include/barry/models/geese/geese-meat.hpp	182
include/barry/models/geese/geese-node-bones hop	182

## **Module Documentation**

### 5.1 Counting

#### **Classes**

- · class NetworkData
  - Data class for Networks.
- · class NodeData

Data definition for the PhyloArray class.

class Counter< Array\_Type, Data\_Type >

A counter function based on change statistics.

#### 5.1.1 Detailed Description

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

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

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

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

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

#### 5.2 Statistical Models

Statistical models available in barry.

#### **Classes**

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

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

· class Flock

A Flock is a group of Geese.

· class Geese

Annotated Phylo Model.

#### 5.2.1 Detailed Description

Statistical models available in barry.

#### 5.3 Network counters

Counters for network models.

#### **Functions**

• void counter\_edges (NetCounters \*counters)

Number of edges.

void counter isolates (NetCounters \*counters)

Number of isolated vertices.

void counter\_mutual (NetCounters \*counters)

Number of mutual ties.

- void counter\_istar2 (NetCounters \*counters)
- void counter\_ostar2 (NetCounters \*counters)
- void counter ttriads (NetCounters \*counters)
- void counter ctriads (NetCounters \*counters)
- void counter\_density (NetCounters \*counters)
- void counter\_idegree15 (NetCounters \*counters)
- void counter\_odegree15 (NetCounters \*counters)
- void counter\_absdiff (NetCounters \*counters, uint attr\_id, double alpha=1.0)

Sum of absolute attribute difference between ego and alter.

• void counter\_diff (NetCounters \*counters, uint attr\_id, double alpha=1.0, double tail\_head=true)

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

- NETWORK\_COUNTER (init\_single\_attr)
- void counter nodeicov (NetCounters \*counters, uint attr id)
- void counter\_nodeocov (NetCounters \*counters, uint attr\_id)
- void counter\_nodecov (NetCounters \*counters, uint attr\_id)
- void counter\_nodematch (NetCounters \*counters, uint attr\_id)
- void counter\_idegree (NetCounters \*counters, std::vector< uint > d)

Counts number of vertices with a given in-degree.

void counter\_odegree (NetCounters \*counters, std::vector< uint > d)

Counts number of vertices with a given out-degree.

void counter degree (NetCounters \*counters, std::vector< uint > d)

Counts number of vertices with a given out-degree.

#### 5.3.1 Detailed Description

Counters for network models.

5.3 Network counters

#### **Parameters**

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

#### 5.3.2 Function Documentation

#### 5.3.2.1 counter\_absdiff()

Sum of absolute attribute difference between ego and alter.

Definition at line 419 of file network.hpp.

#### 5.3.2.2 counter\_ctriads()

Definition at line 322 of file network.hpp.

#### 5.3.2.3 counter\_degree()

Counts number of vertices with a given out-degree.

Definition at line 690 of file network.hpp.

#### 5.3.2.4 counter\_density()

Definition at line 361 of file network.hpp.

#### 5.3.2.5 counter\_diff()

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

Definition at line 461 of file network.hpp.

#### 5.3.2.6 counter\_edges()

Number of edges.

Definition at line 128 of file network.hpp.

#### 5.3.2.7 counter\_idegree()

Counts number of vertices with a given in-degree.

Definition at line 604 of file network.hpp.

#### 5.3.2.8 counter\_idegree15()

Definition at line 377 of file network.hpp.

#### 5.3.2.9 counter\_isolates()

Number of isolated vertices.

Definition at line 142 of file network.hpp.

5.3 Network counters

#### 5.3.2.10 counter\_istar2()

Definition at line 210 of file network.hpp.

#### 5.3.2.11 counter\_mutual()

Number of mutual ties.

Definition at line 172 of file network.hpp.

#### 5.3.2.12 counter\_nodecov()

Definition at line 558 of file network.hpp.

#### 5.3.2.13 counter\_nodeicov()

Definition at line 520 of file network.hpp.

#### 5.3.2.14 counter\_nodematch()

Definition at line 578 of file network.hpp.

#### 5.3.2.15 counter\_nodeocov()

Definition at line 539 of file network.hpp.

#### 5.3.2.16 counter\_odegree()

Counts number of vertices with a given out-degree.

Definition at line 646 of file network.hpp.

#### 5.3.2.17 counter\_odegree15()

Definition at line 397 of file network.hpp.

#### 5.3.2.18 counter\_ostar2()

Definition at line 228 of file network.hpp.

#### 5.3.2.19 counter\_ttriads()

Definition at line 247 of file network.hpp.

5.4 Phylo counters 15

#### 5.3.2.20 NETWORK\_COUNTER()

Definition at line 503 of file network.hpp.

### 5.4 Phylo counters

Counters for phylogenetic modeling.

#### **Functions**

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

k genes gain function nfun

void counter\_genes\_changing (PhyloCounters \*counters, bool duplication=true)

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

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

Overall functional loss.

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

Total count of losses for an specific function.

- void counter\_overall\_changes (PhyloCounters \*counters, bool duplication=true)
  - Total number of changes. Use this statistic to account for "preservation".
- void counter\_subfun (PhyloCounters \*counters, uint nfunA, uint nfunB, bool duplication=true)

Total count of Sub-functionalization events.

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

Co-evolution (joint gain or loss)

- void counter\_longest (PhyloCounters \*counters)
  - Longest branch mutates (either by gain or by loss)
- void counter\_neofun (PhyloCounters \*counters, uint nfunA, uint nfunB, bool duplication=true)

Total number of neofunctionalization events.

• void counter\_neofun\_a2b (PhyloCounters \*counters, uint nfunA, uint nfunB, bool duplication=true)

Total number of neofunctionalization events.

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

#### 5.4.1 Detailed Description

Counters for phylogenetic modeling.

#### **Parameters**

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

#### 5.4.2 Function Documentation

#### 5.4.2.1 counter\_co\_opt()

Function co-opting.

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

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

This algorithm implements the change statistic.

Definition at line 1084 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 714 of file phylo.hpp.

5.4 Phylo counters

#### 5.4.2.3 counter\_gains()

Functional gains for a specific function (nfun).

Definition at line 152 of file phylo.hpp.

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

k genes gain function nfun

Definition at line 194 of file phylo.hpp.

#### 5.4.2.5 counter\_genes\_changing()

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

Definition at line 268 of file phylo.hpp.

#### 5.4.2.6 counter\_longest()

Longest branch mutates (either by gain or by loss)

Definition at line 773 of file phylo.hpp.

18 Module Documentation

#### 5.4.2.7 counter\_loss()

Total count of losses for an specific function.

Definition at line 515 of file phylo.hpp.

#### 5.4.2.8 counter\_maxfuns()

Cap the number of functions per gene.

Definition at line 431 of file phylo.hpp.

#### 5.4.2.9 counter\_neofun()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 878 of file phylo.hpp.

#### 5.4.2.10 counter\_neofun\_a2b()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 963 of file phylo.hpp.

5.4 Phylo counters

#### 5.4.2.11 counter\_overall\_changes()

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

Definition at line 562 of file phylo.hpp.

#### 5.4.2.12 counter\_overall\_gains()

Overall functional gains.

Total number of gains (irrespective of the function).

Definition at line 112 of file phylo.hpp.

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

Overall functional loss.

Definition at line 385 of file phylo.hpp.

#### 5.4.2.14 counter\_subfun()

Total count of Sub-functionalization events.

It requires to specify data = {funA, funB}

Definition at line 628 of file phylo.hpp.

20 Module Documentation

## 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, bool duplication=true) 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()

```
void rule_dyn_limit_changes (
          PhyloSupport * support,
          uint pos,
          uint lb,
          uint ub,
          bool duplication = true ) [inline]
```

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 21

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

Definition at line 1217 of file phylo.hpp.

22 Module Documentation

# **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 20 of file typedefs.hpp.

#### 6.5.2.2 NONE

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

Definition at line 21 of file typedefs.hpp.

#### 6.5.2.3 ONE

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

Definition at line 22 of file typedefs.hpp.

## 6.5.2.4 TWO

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

Definition at line 23 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 38 of file typedefs.hpp.

#### 6.6.2.2 AS\_ZERO

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

Definition at line 37 of file typedefs.hpp.

#### 6.6.2.3 BOTH

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

Definition at line 31 of file typedefs.hpp.

#### 6.6.2.4 NONE

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

Definition at line 32 of file typedefs.hpp.

#### 6.6.2.5 ONE

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

Definition at line 33 of file typedefs.hpp.

#### 6.6.2.6 TWO

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

Definition at line 34 of file typedefs.hpp.

#### 6.6.2.7 UKNOWN

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

Definition at line 36 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
- const Row\_type< Cell\_Type > \* get\_row (uint i, bool check\_bounds=true) const
- const Col type< Cell Type > \* get col (uint i, bool check bounds=true) const
- std::vector< Cell\_Type > get\_col\_vec (uint i, bool check\_bounds=true) const
- std::vector< Cell\_Type > get\_row\_vec (uint i, bool check\_bounds=true) const
- void get\_col\_vec (std::vector< Cell\_Type > \*x, uint i, bool check\_bounds=true) const
- void get\_row\_vec (std::vector< Cell\_Type > \*x, uint i, bool check\_bounds=true) const
- const Row\_type< Cell\_Type > & row (uint i, bool check\_bounds=true) const
- const Col\_type< Cell\_Type > & col (uint i, bool check\_bounds=true) const
- Entries < Cell\_Type > get\_entries () const

#### Get the edgelist.

- void transpose ()
- void clear (bool hard=true)
- void resize (uint N\_, uint M\_)
- void reserve ()
- void print () const

#### Constructors

#### **Parameters**

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

• BArray ()

Zero-size array.

BArray (uint N\_, uint M\_)

Empty array.

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

Edgelist with data.

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

Edgelist with no data (simpler)

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

Move operator.

- BArray< Cell\_Type, Data\_Type > & operator= (BArray< Cell\_Type, Data\_Type > &&x) noexcept
   Move assignment.
- 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.

- BArray< Cell\_Type, Data\_Type > & operator+= (const std::pair< uint, uint > &coords)
- BArray< Cell\_Type, Data\_Type > & operator-= (const std::pair< uint, uint > &coords)

- BArrayCell< Cell\_Type, Data\_Type > operator() (uint i, uint j, bool check\_bounds=true)
- const BArrayCell\_const< Cell\_Type, Data\_Type > operator() (uint i, uint j, bool check\_bounds=true) const
- void rm\_cell (uint i, uint j, bool check\_bounds=true, bool check\_exists=true)
- void insert\_cell (uint i, uint j, const Cell< Cell\_Type > &v, bool check\_bounds, bool check\_exists)
- void insert\_cell (uint i, uint j, Cell
   Cell\_Type > &&v, bool check\_bounds, bool check\_exists)
- void insert\_cell (uint i, uint j, Cell\_Type v, bool check\_bounds, bool check\_exists)
- void swap\_cells (uint i0, uint j0, uint i1, uint j1, bool check\_bounds=true, int check\_exists=CHECK::BOTH, int \*report=nullptr)
- void toggle\_cell (uint i, uint j, bool check\_bounds=true, int check\_exists=EXISTS::UKNOWN)
- void toggle lock (uint i, uint j, bool check bounds=true)

#### Column/row wise interchange

- void swap\_rows (uint i0, uint i1, bool check\_bounds=true)
- void swap\_cols (uint j0, uint j1, bool check\_bounds=true)
- void zero row (uint i, bool check bounds=true)
- void zero\_col (uint j, bool check\_bounds=true)

#### **Arithmetic operators**

```
• BArray< Cell_Type, Data_Type > & operator+= (const BArray< Cell_Type, Data_Type > &rhs)
```

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

#### **Public Attributes**

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 60 of file barray-bones.hpp.

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

Empty array.

Definition at line 63 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()

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

#### 7.1.3.3 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 get\_cell()

#### 7.1.3.7 get\_col()

#### 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()

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

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

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

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

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

#### 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()

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

#### 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

## ${\bf 7.1.4.1 \quad BArrayCell} < {\bf Cell\_Type, Data\_Type} >$

```
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 45 of file barray-bones.hpp.

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

• include/barry/barray-bones.hpp

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

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

#### **Public Member Functions**

- BArrayCell (BArray < Cell\_Type, Data\_Type > \*Array\_, uint i\_, uint j\_, bool check\_bounds=true)
- ∼BArrayCell ()
- void operator= (const Cell\_Type &val)
- void operator+= (const Cell\_Type &val)
- void operator-= (const Cell\_Type &val)
- void operator\*= (const Cell\_Type &val)
- void operator/= (const Cell\_Type &val)
- operator Cell\_Type () const
- bool operator== (const Cell\_Type &val) const

#### 7.2.1 Detailed Description

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

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

#### 7.2.2 Constructor & Destructor Documentation

#### 7.2.2.1 BArrayCell()

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

#### 7.2.2.2 ∼BArrayCell()

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

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

#### 7.2.3 Member Function Documentation

#### 7.2.3.1 operator Cell\_Type()

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

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

#### 7.2.3.2 operator\*=()

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

#### 7.2.3.3 operator+=()

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

#### 7.2.3.4 operator-=()

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

#### 7.2.3.5 operator/=()

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

#### 7.2.3.6 operator=()

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

#### 7.2.3.7 operator==()

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

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

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

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

#include <barraycell-bones.hpp>

#### **Public Member Functions**

- BArrayCell\_const (const BArray < Cell\_Type, Data\_Type > \*Array\_, uint i\_, uint j\_, bool check\_bounds=true)
- ∼BArrayCell const ()
- operator Cell\_Type () const
- bool operator== (const Cell\_Type &val) const
- bool operator!= (const Cell\_Type &val) const
- bool operator< (const Cell\_Type &val) const
- bool operator> (const Cell\_Type &val) const
- bool operator<= (const Cell\_Type &val) const
- bool operator>= (const Cell\_Type &val) const

#### 7.3.1 Detailed Description

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

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

#### 7.3.2 Constructor & Destructor Documentation

#### 7.3.2.1 BArrayCell const()

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

#### 7.3.2.2 ~BArrayCell\_const()

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

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

#### 7.3.3 Member Function Documentation

#### 7.3.3.1 operator Cell\_Type()

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

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

#### 7.3.3.2 operator"!=()

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

#### 7.3.3.3 operator<()

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

#### 7.3.3.4 operator<=()

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

#### 7.3.3.5 operator==()

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

#### 7.3.3.6 operator>()

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

#### 7.3.3.7 operator>=()

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

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

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

## 7.4 BArrayCol < Cell\_Type, Data\_Type > Class Template Reference

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

#### **Public Member Functions**

```
    BArrayCol (BArray < Cell_Type, Data_Type > *Array_, uint i_, bool check_bounds=true)
```

- ∼BArrayCol ()
- 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
- Col\_type< Cell\_Type >::iterator begin () noexcept
- Col\_type< Cell\_Type >::iterator end () noexcept

#### 7.4.1 Detailed Description

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

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

## 7.4.2 Constructor & Destructor Documentation

## 7.4.2.1 BArrayCol()

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

#### 7.4.2.2 ~BArrayCol()

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

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

#### 7.4.3 Member Function Documentation

## 7.4.3.1 begin()

```
template<typename Cell_Type , typename Data_Type >
Col_type< Cell_Type >::iterator BArrayCol< Cell_Type, Data_Type >::begin [inline], [noexcept]
```

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

#### 7.4.3.2 end()

```
template<typename Cell_Type , typename Data_Type >
Col_type< Cell_Type >::iterator BArrayCol< Cell_Type, Data_Type >::end [inline], [noexcept]
```

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

#### 7.4.3.3 operator Cell\_Type()

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

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

#### 7.4.3.4 operator\*=()

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

#### 7.4.3.5 operator+=()

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

#### 7.4.3.6 operator-=()

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

#### 7.4.3.7 operator/=()

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

#### 7.4.3.8 operator=()

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

#### 7.4.3.9 operator==()

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

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

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

# 7.5 BArrayCol\_const< Cell\_Type, Data\_Type > Class Template Reference

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

#### **Public Member Functions**

- BArrayCol (const BArray < Cell\_Type, Data\_Type > \*Array\_, uint i\_, bool check\_bounds=true)
- ∼BArrayCol\_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

#### 7.5.1 Detailed Description

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

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

#### 7.5.2 Constructor & Destructor Documentation

## 7.5.2.1 $\sim$ BArrayCol\_const()

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

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

#### 7.5.3 Member Function Documentation

#### 7.5.3.1 BArrayCol()

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

## 7.5.3.2 operator"!=()

Definition at line 92 of file barraycol-meat.hpp.

#### 7.5.3.3 operator<()

Definition at line 97 of file barraycol-meat.hpp.

#### 7.5.3.4 operator<=()

Definition at line 107 of file barraycol-meat.hpp.

#### 7.5.3.5 operator==()

Definition at line 87 of file barraycol-meat.hpp.

#### 7.5.3.6 operator>()

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

#### 7.5.3.7 operator>=()

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

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

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

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

Dense bi-dimensional array.

#include <barraydense-bones.hpp>

#### **Public Member Functions**

```
• BArrayDense ()
```

- BArrayDense (uint N\_, uint M\_, std::vector< Cell\_Type > elements\_={})
- ∼BArrayDense ()
- void fill (const Cell\_Type &d)
- const std::vector< Cell\_Type > & elements\_raw () const noexcept
- const std::vector< Cell\_Type > \* elements\_ptr () const noexcept
- uint nrow () const noexcept
- uint ncol () const noexcept
- Cell\_Type operator() (uint i, uint j, bool check\_bounds=true) const
- Cell\_Type operator[] (uint i) const
- · void print () const

#### 7.6.1 Detailed Description

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

Dense bi-dimensional array.

elements is stored in a std::vector, in col-major order.

**Template Parameters** 

```
Cell_Type
```

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

#### 7.6.2 Constructor & Destructor Documentation

#### 7.6.2.1 BArrayDense() [1/2]

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

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

#### 7.6.2.2 BArrayDense() [2/2]

```
template<typename Cell_Type , typename Data_Type >
BArrayDense< Cell_Type, Data_Type >::BArrayDense (
          uint N_,
          uint M_,
          std::vector< Cell_Type > elements_ = {} ) [inline]
```

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

#### 7.6.2.3 ∼BArrayDense()

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

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

#### 7.6.3 Member Function Documentation

#### 7.6.3.1 elements\_ptr()

```
template<typename Cell_Type , typename Data_Type >
const std::vector< Cell_Type > * BArrayDense< Cell_Type, Data_Type >::elements_ptr [inline],
[noexcept]
```

Definition at line 85 of file barraydense-bones.hpp.

#### 7.6.3.2 elements\_raw()

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

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

#### 7.6.3.3 fill()

Definition at line 70 of file barraydense-bones.hpp.

#### 7.6.3.4 ncol()

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

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

#### 7.6.3.5 nrow()

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

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

#### 7.6.3.6 operator()()

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

#### 7.6.3.7 operator[]()

Definition at line 122 of file barraydense-bones.hpp.

#### 7.6.3.8 print()

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

Definition at line 128 of file barraydense-bones.hpp.

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

• include/barry/barraydense-bones.hpp

# 7.7 Cell< Cell\_Type > Class Template Reference

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

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

# **Public Member Functions**

- Cell ()
- Cell (Cell\_Type value\_, bool visited\_=false)
- ~Cell ()
- Cell (const Cell< Cell\_Type > &arg)
- Cell< Cell\_Type > & operator= (Cell< Cell\_Type > &other)
- Cell (Cell< Cell\_Type > &&arg) noexcept
- Cell< Cell\_Type > & operator= (Cell< Cell\_Type > &&other) noexcept
- void add (Cell Type x)
- operator Cell\_Type () const
- void add (double x)
- void add (unsigned int x)
- void add (int x)
- Cell ()
- Cell ()
- Cell ()

#### **Public Attributes**

- Cell\_Type value
- bool visited

# 7.7.1 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.

#### 7.7.2 Constructor & Destructor Documentation

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

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

# 7.7.2.2 Cell() [2/7]

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

#### 7.7.2.3 ∼CeII()

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

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

# 7.7.2.4 Cell() [3/7]

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

# 7.7.2.5 Cell() [4/7]

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

#### 7.7.2.6 Cell() [5/7]

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

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

# 7.7.2.7 Cell() [6/7]

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

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

#### 7.7.2.8 Cell() [7/7]

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

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

#### 7.7.3 Member Function Documentation

# 7.7.3.1 add() [1/4]

#### 7.7.3.2 add() [2/4]

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

# 7.7.3.3 add() [3/4]

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

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

#### 7.7.3.4 add() [4/4]

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

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

# 7.7.3.5 operator Cell\_Type()

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

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

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

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

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

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

# 7.7.4 Member Data Documentation

#### 7.7.4.1 value

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

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

#### 7.7.4.2 visited

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

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

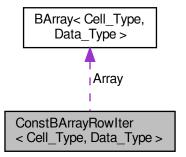
The documentation for this class was generated from the following files:

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

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

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

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



#### **Public Member Functions**

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

#### **Public Attributes**

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

# 7.8.1 Detailed Description

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

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

#### 7.8.2 Constructor & Destructor Documentation

#### 7.8.2.1 ConstBArrayRowIter()

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

#### 7.8.2.2 ∼ConstBArrayRowlter()

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

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

#### 7.8.3 Member Data Documentation

#### 7.8.3.1 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.

#### 7.8.3.2 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.

#### 7.8.3.3 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.

#### 7.8.3.4 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

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

A counter function based on change statistics.

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

# **Public Member Functions**

- $\sim$ Counter ()
- double count (Array\_Type &Array, uint i, uint j)
- double init (Array\_Type &Array, uint i, uint j)

#### Creator passing a counter and an initializer

#### **Parameters**

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

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

- Counter (const Counter < Array\_Type, Data\_Type > &counter\_)
   Copy constructor.
- $\bullet \ \, {\tt Counter} ({\tt Counter} < {\tt Array\_Type}, \, {\tt Data\_Type} > \& {\tt counter\_)} \, \, {\tt noexcept} \\$

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 = nullptr
- bool delete data = false
- std::string name = ""
- std::string desc = ""

#### 7.9.1 Detailed Description

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

A counter function based on change statistics.

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

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

#### 7.9.2 Constructor & Destructor Documentation

# 7.9.2.1 Counter() [1/4]

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

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

#### 7.9.2.2 Counter() [2/4]

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

#### 7.9.2.3 Counter() [3/4]

Copy constructor.

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

#### 7.9.2.4 Counter() [4/4]

Move constructor.

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

# 7.9.2.5 ~Counter()

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

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

#### 7.9.3 Member Function Documentation

#### 7.9.3.1 count()

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

#### 7.9.3.2 init()

Definition at line 126 of file counters-meat.hpp.

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

Copy assignment.

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

# 7.9.3.4 operator=() [2/2]

Move assignment.

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

#### 7.9.4 Member Data Documentation

#### 7.9.4.1 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.

#### 7.9.4.2 data

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

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

#### 7.9.4.3 delete\_data

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

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

#### 7.9.4.4 desc

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

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

#### 7.9.4.5 init fun

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

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

#### 7.9.4.6 name

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

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

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

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

# 7.10 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< Array\_Type, Data\_Type > \*counter)
- void add\_counter (Counter\_fun\_type< Array\_Type, Data\_Type > count\_fun\_, Counter\_fun\_type< Array\_
   —
   Type, Data\_Type > init\_fun\_=nullptr, Data\_Type \*data\_=nullptr, bool delete\_data\_=false, std::string name
   —="", std::string desc\_="")
- void clear ()

#### 7.10.1 Detailed Description

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

Vector of counters.

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

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

#### 7.10.2 Constructor & Destructor Documentation

#### 7.10.2.1 Counters() [1/3]

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

Definition at line 143 of file counters-meat.hpp.

#### 7.10.2.2 ~Counters()

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

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

#### 7.10.2.3 Counters() [2/3]

Copy constructor.

**Parameters** 



Definition at line 160 of file counters-meat.hpp.

# **7.10.2.4 Counters()** [3/3]

Move constructor.

#### **Parameters**



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

#### 7.10.3 Member Function Documentation

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

Definition at line 276 of file counters-meat.hpp.

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

Definition at line 288 of file counters-meat.hpp.

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

Definition at line 299 of file counters-meat.hpp.

#### 7.10.3.4 clear()

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

Definition at line 328 of file counters-meat.hpp.

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

Copy assignment constructor.

#### **Parameters**



#### Returns

Counters<Array\_Type,Data\_Type>

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

# 7.10.3.6 operator=() [2/2]

Move assignment constructor.

#### **Parameters**



#### Returns

 ${\tt Counters}{<}{\tt Array\_Type}, {\tt Data\_Type}{>} \&$ 

Definition at line 248 of file counters-meat.hpp.

#### 7.10.3.7 operator[]()

Returns a pointer to a particular counter.

#### **Parameters**

```
idx Id of the counter
```

#### Returns

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

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

#### 7.10.3.8 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 159 of file counters-bones.hpp.

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

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

# 7.11 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

# 7.11.1 Detailed Description

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

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

**Template Parameters** 

```
Cell_Type Any type
```

Definition at line 59 of file typedefs.hpp.

# 7.11.2 Constructor & Destructor Documentation

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

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

Definition at line 65 of file typedefs.hpp.

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

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

Definition at line 66 of file typedefs.hpp.

#### 7.11.2.3 ∼Entries()

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

Definition at line 73 of file typedefs.hpp.

#### 7.11.3 Member Function Documentation

# 7.11.3.1 resize()

Definition at line 75 of file typedefs.hpp.

#### 7.11.4 Member Data Documentation

#### 7.11.4.1 source

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

Definition at line 61 of file typedefs.hpp.

#### 7.11.4.2 target

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

Definition at line 62 of file typedefs.hpp.

#### 7.11.4.3 val

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

Definition at line 63 of file typedefs.hpp.

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

• include/barry/typedefs.hpp

7.12 Flock Class Reference 73

#### 7.12 Flock Class Reference

A Flock is a group of Geese.

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

#### **Public Member Functions**

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

Add a tree to the flock.

• void set\_seed (const unsigned int &s)

Set the seed of the model.

- void init ()
- phylocounters::PhyloCounters \* get\_counters ()
- phylocounters::PhyloSupport \* get\_support ()

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

# **Public Attributes**

- std::vector< Geese > dat
- unsigned int nfunctions = 0u
- bool initialized = false
- std::mt19937 rengine
- phylocounters::PhyloModel support = phylocounters::PhyloModel()

#### 7.12.1 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.

# 7.12.2 Constructor & Destructor Documentation

# 7.12.2.1 Flock()

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

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

# 7.12.2.2 ∼Flock()

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

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

# 7.12.3 Member Function Documentation

# 7.12.3.1 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-meet.hpp.

7.12 Flock Class Reference 75

# 7.12.3.2 get\_counters()

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

Definition at line 69 of file flock-meet.hpp.

#### 7.12.3.3 get\_support()

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

Definition at line 78 of file flock-meet.hpp.

# 7.12.3.4 init()

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

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

# 7.12.3.5 likelihood\_joint()

Returns the joint likelihood of the model.

#### **Parameters**

par	Vector of model parameters.
as_log	When true it will return the value as log.
use_reduced_sequence	When true (default) will compute the likelihood using the reduced sequence, which is faster.

#### Returns

double

Definition at line 82 of file flock-meet.hpp.

#### 7.12.3.6 nfuns()

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

Definition at line 107 of file flock-meet.hpp.

# 7.12.3.7 nleafs()

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

Definition at line 130 of file flock-meet.hpp.

#### 7.12.3.8 nnodes()

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

Definition at line 119 of file flock-meet.hpp.

# 7.12.3.9 nterms()

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

Definition at line 142 of file flock-meet.hpp.

#### 7.12.3.10 ntrees()

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

Definition at line 113 of file flock-meet.hpp.

# 7.12.3.11 operator()()

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

Access the i-th geese element.

7.12 Flock Class Reference 77

#### **Parameters**

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

#### Returns

Geese \*

Definition at line 155 of file flock-meet.hpp.

# 7.12.3.12 set\_seed()

Set the seed of the model.

#### **Parameters**

```
s Passed to the rengine.seed() member object.
```

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

### 7.12.3.13 support\_size()

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

Definition at line 149 of file flock-meet.hpp.

# 7.12.4 Member Data Documentation

# 7.12.4.1 dat

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

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

#### 7.12.4.2 initialized

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

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

#### 7.12.4.3 nfunctions

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

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

#### 7.12.4.4 rengine

```
std::mt19937 Flock::rengine
```

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

#### 7.12.4.5 support

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

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

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

- include/barry/models/geese/flock-bones.hpp
- include/barry/models/geese/flock-meet.hpp

# 7.13 FreqTable < T > Class Template Reference

Database of statistics.

```
#include <statsdb.hpp>
```

# **Public Member Functions**

- FreqTable ()
- ∼FreqTable ()
- void add (const std::vector< T > &x)
- Counts\_type as\_vector () const
- MapVec\_type< T, uint > get\_data () const
- const MapVec\_type< T, uint > \* get\_data\_ptr () const
- void clear ()
- void reserve (unsigned int n)
- void print () const
- size\_t size () const noexcept

# 7.13.1 Detailed Description

```
template<typename T = double> class FreqTable< T >
```

Database of statistics.

This is mostly used in Support.

Definition at line 16 of file statsdb.hpp.

#### 7.13.2 Constructor & Destructor Documentation

#### 7.13.2.1 FreqTable()

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

Definition at line 28 of file statsdb.hpp.

# 7.13.2.2 $\sim$ FreqTable()

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

Definition at line 29 of file statsdb.hpp.

# 7.13.3 Member Function Documentation

# 7.13.3.1 add()

Definition at line 47 of file statsdb.hpp.

# 7.13.3.2 as\_vector()

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

Definition at line 61 of file statsdb.hpp.

# 7.13.3.3 clear()

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

Definition at line 83 of file statsdb.hpp.

#### 7.13.3.4 get\_data()

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

Definition at line 73 of file statsdb.hpp.

# 7.13.3.5 get\_data\_ptr()

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

Definition at line 78 of file statsdb.hpp.

#### 7.13.3.6 print()

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

Definition at line 102 of file statsdb.hpp.

#### 7.13.3.7 reserve()

Definition at line 89 of file statsdb.hpp.

#### 7.13.3.8 size()

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

Definition at line 126 of file statsdb.hpp.

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

include/barry/statsdb.hpp

#### 7.14 Geese Class Reference

Annotated Phylo Model.

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

#### **Public Member Functions**

- ~Geese ()
- void init ()
- void inherit\_support (const Geese &model\_, bool delete\_support\_=false)
- void calc\_sequence (Node \*n=nullptr)
- void calc\_reduced\_sequence ()
- double likelihood (const std::vector< double > &par, bool as log=false, bool use reduced sequence=true)
- double likelihood\_exhaust (const std::vector< double > &par)
- std::vector< double > get\_probabilities () const
- void set\_seed (const unsigned int &s)
- std::vector< std::vector< unsigned int > > simulate (const std::vector< double > &par)
- std::vector< std::vector< double >> observed counts ()
- void print\_observed\_counts ()
- void init\_node (Node &n)
- void update\_annotations (unsigned int nodeid, std::vector< unsigned int > newann)
- std::mt19937 \* get\_rengine ()
- phylocounters::PhyloCounters \* get\_counters ()
- phylocounters::PhyloSupport \* get support ()
- $std::vector < std::vector < bool >> get\_states ()$

#### 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 $\mathbb{N}$ .
parent	Id of the parent gene. Also of length ${\tt N}$

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

#### Information about the model

- · unsigned int nfuns () const noexcept
- unsigned int nnodes () const noexcept
- unsigned int nleafs () const noexcept
- unsigned int nterms () const
- unsigned int support\_size () const noexcept

#### Geese prediction

Calculate the conditional probability

#### **Parameters**

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

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

#### Returns

std::vector< double > Returns the posterior probability

- std::vector< std::vector< double >> predict (const std::vector< double > &par, std::vector< std::vector< double >> \*res\_prob=nullptr, bool leave\_one\_out=false, bool only\_annotated=false, bool use\_reduced -\_sequence=true)
- std::vector < std::vector < double > > predict\_backend (const std::vector < double > &par, bool use\_←
  reduced\_sequence, const std::vector < uint > &preorder)

# **Public Attributes**

unsigned int nfunctions

- std::map< unsigned int, Node > nodes
- barry::MapVec\_type< unsigned int > map\_to\_nodes
- std::vector< unsigned int > sequence
- std::vector< unsigned int > reduced\_sequence
- bool initialized = false
- bool delete\_rengine = false
- bool delete\_support = false

# 7.14.1 Detailed Description

Annotated Phylo Model.

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

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

#### 7.14.2 Constructor & Destructor Documentation

#### 7.14.2.1 Geese() [1/4]

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

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

# 7.14.2.2 Geese() [2/4]

```
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.

# 7.14.2.3 Geese() [3/4]

Definition at line 163 of file geese-meat-constructors.hpp.

#### 7.14.2.4 Geese() [4/4]

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

# 7.14.2.5 ∼Geese()

```
Geese::∼Geese ( ) [inline]
```

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

#### 7.14.3 Member Function Documentation

# 7.14.3.1 calc\_reduced\_sequence()

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

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

# 7.14.3.2 calc\_sequence()

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

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

# 7.14.3.3 get\_counters()

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

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

#### 7.14.3.4 get\_probabilities()

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

Definition at line 267 of file geese-meat.hpp.

# 7.14.3.5 get\_rengine()

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

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

# 7.14.3.6 get\_states()

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

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

# 7.14.3.7 get\_support()

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

Definition at line 435 of file geese-meat.hpp.

# 7.14.3.8 inherit\_support()

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

# 7.14.3.9 init()

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

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

#### 7.14.3.10 init\_node()

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

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

#### 7.14.3.11 likelihood()

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

#### 7.14.3.12 likelihood\_exhaust()

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

#### 7.14.3.13 nfuns()

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

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

#### 7.14.3.14 nleafs()

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

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

# 7.14.3.15 nnodes()

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

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

#### 7.14.3.16 nterms()

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

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

#### 7.14.3.17 observed\_counts()

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

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

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

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

# 7.14.3.20 predict()

Definition at line 166 of file geese-meat-predict.hpp.

#### 7.14.3.21 predict\_backend()

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

#### 7.14.3.22 print\_observed\_counts()

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

Definition at line 365 of file geese-meat.hpp.

#### 7.14.3.23 set\_seed()

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

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

# 7.14.3.24 simulate()

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

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

# 7.14.3.25 support\_size()

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

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

# 7.14.3.26 update\_annotations()

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

Definition at line 158 of file geese-meat.hpp.

# 7.14.4 Member Data Documentation

## 7.14.4.1 delete\_rengine

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

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

#### 7.14.4.2 delete\_support

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

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

#### 7.14.4.3 initialized

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

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

## 7.14.4.4 map\_to\_nodes

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

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

#### **7.14.4.5** nfunctions

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

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

#### 7.14.4.6 nodes

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

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

#### 7.14.4.7 reduced\_sequence

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

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

#### 7.14.4.8 sequence

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

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

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

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

# 7.15 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\_)
- uint add\_array (const Array\_Type &Array\_, bool force\_new=false)

Adds an array to the support of not already included.

- void print\_stats (uint i) const
- Array\_Type sample (const Array\_Type &Array\_, const std::vector< double > &params={})
- Array\_Type sample (const uint &i, const std::vector< double > &params)

- 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 ()

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

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

- void add\_counter (Counter< Array\_Type, Data\_Counter\_Type > &counter)
- void add\_counter (Counter< Array\_Type, Data\_Counter\_Type > \*counter)
- void add\_counter (Counter\_fun\_type < Array\_Type, Data\_Counter\_Type > count\_fun\_, Counter\_fun\_type <
   Array\_Type, Data\_Counter\_Type > init\_fun\_=nullptr, Data\_Counter\_Type \*data\_=nullptr, bool delete\_
   data =false)
- void set\_counters (Counters < Array\_Type, Data\_Counter\_Type > \*counters\_)

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

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

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

## Likelihood functions.

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

#### **Parameters**

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

- double likelihood (const std::vector< double > &params, const uint &i, bool as\_log=false)
- double likelihood (const std::vector< double > &params, const Array\_Type &Array\_, int i=-1, bool as\_←
  log=false)
- double likelihood (const std::vector< double > &params, const std::vector< double > &target\_, const uint &i, bool as log=false)
- double likelihood total (const std::vector< double > &params, bool as log=false)

## Extract elements by index

#### Parameters

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

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

#### Size of the model

Number of different supports included in the model

This will return the size of stats.

#### Returns

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

- unsigned int size () const noexcept
- unsigned int size\_unique () const noexcept
- · unsigned int nterms () const noexcept
- unsigned int support size () const noexcept

## 7.15.1 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 104 of file model-bones.hpp.

#### 7.15.2 Constructor & Destructor Documentation

## 7.15.2.1 Model() [1/3]

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

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

#### 7.15.2.2 Model() [2/3]

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

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

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

#### 7.15.2.4 ~Model()

```
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 ( ) [inline]
```

Definition at line 204 of file model-bones.hpp.

#### 7.15.3 Member Function Documentation

#### 7.15.3.1 add\_array()

Adds an array to the support of not already included.

#### **Parameters**

Array_	array to be added
force_new	If false, it will use keygen to obtain a double vector and create a hash of it. If the hash has
	been computed earlier, the support is recycled.

#### Returns

The number of the array.

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

Definition at line 333 of file model-meat.hpp.

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

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

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

Definition at line 176 of file model-meat.hpp.

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

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

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

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

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

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

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

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

## 7.15.3.8 add\_rule\_dyn() [1/3]

Definition at line 279 of file model-meat.hpp.

#### 7.15.3.9 add\_rule\_dyn() [2/3]

Definition at line 288 of file model-meat.hpp.

## 7.15.3.10 add\_rule\_dyn() [3/3]

Definition at line 298 of file model-meat.hpp.

#### 7.15.3.11 get\_counters()

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

Counters< Array_Type, Data_Counter_Type > * Model< Array_Type, Data_Counter_Type, Data_Rule

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

Definition at line 713 of file model-meat.hpp.

#### 7.15.3.12 get\_norm\_const()

Definition at line 567 of file model-meat.hpp.

#### 7.15.3.13 get pset()

Definition at line 599 of file model-meat.hpp.

## 7.15.3.14 get\_rengine()

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

Definition at line 707 of file model-meat.hpp.

#### 7.15.3.15 get\_rules()

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

Definition at line 719 of file model-meat.hpp.

## 7.15.3.16 get\_rules\_dyn()

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

Definition at line 725 of file model-meat.hpp.

#### 7.15.3.17 get\_stats()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type > const std::vector< std::vector< double > > * Model< Array_Type, Data_Counter_Type, Data_\leftarrow Rule_Type, Data_Rule_Dyn_Type >::get_stats ( const uint & i ) [inline]
```

Definition at line 612 of file model-meat.hpp.

#### 7.15.3.18 get\_support()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > * Model< Array

_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::get_support [inline]
```

Definition at line 731 of file model-meat.hpp.

#### 7.15.3.19 likelihood() [1/3]

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

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

Definition at line 493 of file model-meat.hpp.

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

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

#### 7.15.3.22 likelihood\_total()

Definition at line 527 of file model-meat.hpp.

#### 7.15.3.23 nterms()

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

Definition at line 653 of file model-meat.hpp.

## 7.15.3.24 operator=()

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

## 7.15.3.25 print\_stats()

Definition at line 624 of file model-meat.hpp.

#### 7.15.3.26 sample() [1/2]

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

Definition at line 671 of file model-meat.hpp.

## 7.15.3.28 set\_counters()

Definition at line 205 of file model-meat.hpp.

#### 7.15.3.29 set\_keygen()

Definition at line 159 of file model-meat.hpp.

#### 7.15.3.30 set\_rengine()

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

Definition at line 176 of file model-bones.hpp.

## 7.15.3.31 set\_rules()

Definition at line 261 of file model-meat.hpp.

#### 7.15.3.32 set rules dyn()

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

#### 7.15.3.33 set seed()

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

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

## 7.15.3.34 size()

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

Definition at line 643 of file model-meat.hpp.

#### 7.15.3.35 size\_unique()

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

Definition at line 648 of file model-meat.hpp.

#### 7.15.3.36 store\_psets()

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

Definition at line 151 of file model-meat.hpp.

#### 7.15.3.37 support\_size()

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

Definition at line 660 of file model-meat.hpp.

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

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

# 7.16 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

## 7.16.1 Detailed Description

Data class used to store arbitrary uint or double vectors.

Definition at line 61 of file network.hpp.

#### 7.16.2 Constructor & Destructor Documentation

## 7.16.2.1 NetCounterData() [1/2]

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

Definition at line 67 of file network.hpp.

## 7.16.2.2 NetCounterData() [2/2]

Definition at line 68 of file network.hpp.

#### 7.16.2.3 ∼NetCounterData()

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

Definition at line 73 of file network.hpp.

## 7.16.3 Member Data Documentation

#### 7.16.3.1 indices

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

Definition at line 64 of file network.hpp.

#### 7.16.3.2 numbers

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

Definition at line 65 of file network.hpp.

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

• include/barry/counters/network.hpp

## 7.17 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.

•  $\sim$ NetworkData ()

## **Public Attributes**

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

#### 7.17.1 Detailed Description

Data class for Networks.

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

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

Definition at line 24 of file network.hpp.

#### 7.17.2 Constructor & Destructor Documentation

#### 7.17.2.1 NetworkData() [1/3]

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

Definition at line 30 of file network.hpp.

## 7.17.2.2 NetworkData() [2/3]

Constructor using a single attribute.

#### **Parameters**

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

Definition at line 38 of file network.hpp.

## 7.17.2.3 NetworkData() [3/3]

Constructor using multiple attributes.

#### **Parameters**

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

Definition at line 50 of file network.hpp.

## 7.17.2.4 ~NetworkData()

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

Definition at line 56 of file network.hpp.

#### 7.17.3 Member Data Documentation

#### 7.17.3.1 directed

```
bool NetworkData::directed = true
```

Definition at line 27 of file network.hpp.

## 7.17.3.2 vertex\_attr

```
std::vector< std::vector< double > > NetworkData::vertex_attr
```

Definition at line 28 of file network.hpp.

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

• include/barry/counters/network.hpp

# 7.18 Node Class Reference

A single node for the model.

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

Collaboration diagram for Node:



7.18 Node Class Reference 107

#### **Public Member Functions**

- ∼Node ()
- int get\_parent () const
- bool is\_leaf () const noexcept

#### Construct a new Node object

- Node ()
- Node (unsigned int id\_, unsigned int ord\_, bool duplication\_)
- Node (unsigned int id\_, unsigned int ord\_, std::vector< unsigned int > annotations\_, bool duplication\_)
- Node (Node &&x) noexcept
- Node (const Node &x)

## **Public Attributes**

· unsigned int id

Id of the node (as specified in the input)

· unsigned int ord

Order in which the node was created.

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

Observed annotations (only defined for Geese)

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

Arrays given all possible states.

Node \* parent = nullptr

Parent node.

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

Offspring nodes.

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

ID of the array in the model.

- bool visited = false
- std::vector< double > subtree\_prob

Induced subtree probabilities.

std::vector< double > probability

The probability of observing each state.

## 7.18.1 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.

## 7.18.2 Constructor & Destructor Documentation

#### 7.18.2.1 Node() [1/5]

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

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

#### 7.18.2.2 Node() [2/5]

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

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

#### 7.18.2.3 Node() [3/5]

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

#### 7.18.2.4 Node() [4/5]

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

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

## 7.18.2.5 Node() [5/5]

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

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

7.18 Node Class Reference 109

## 7.18.2.6 ∼Node()

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

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

#### 7.18.3 Member Function Documentation

## 7.18.3.1 get\_parent()

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

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

## 7.18.3.2 is\_leaf()

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

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

#### 7.18.4 Member Data Documentation

#### 7.18.4.1 annotations

```
\verb|std::vector<| unsigned int > \verb|Node::annotations| \\
```

Observed annotations (only defined for Geese)

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

## 7.18.4.2 array

phylocounters::PhyloArray Node::array

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

#### 7.18.4.3 arrays

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

Arrays given all possible states.

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

#### 7.18.4.4 duplication

```
bool Node::duplication
```

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

## 7.18.4.5 id

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

Id of the node (as specified in the input)

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

## 7.18.4.6 narray

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

ID of the array in the model.

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

#### 7.18.4.7 offspring

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

Offspring nodes.

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

7.18 Node Class Reference 111

#### 7.18.4.8 ord

unsigned int Node::ord

Order in which the node was created.

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

#### 7.18.4.9 parent

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

Parent node.

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

## 7.18.4.10 probability

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

The probability of observing each state.

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

## 7.18.4.11 subtree\_prob

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

Induced subtree probabilities.

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

#### 7.18.4.12 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

## 7.19 NodeData Class Reference

Data definition for the PhyloArray class.

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

#### **Public Member Functions**

- NodeData ()
- NodeData (const std::vector< double > &blengths\_, const std::vector< bool > &states\_, bool duplication
   —=true)
- ∼NodeData ()

#### **Public Attributes**

- std::vector< double > blengths
- std::vector< bool > states
- bool duplication = true

## 7.19.1 Detailed Description

Data definition for the PhyloArray class.

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

This holds basic information about a given node.

Definition at line 23 of file phylo.hpp.

#### 7.19.2 Constructor & Destructor Documentation

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

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

Definition at line 41 of file phylo.hpp.

#### 7.19.2.2 NodeData() [2/2]

Definition at line 43 of file phylo.hpp.

#### 7.19.2.3 ~NodeData()

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

Definition at line 49 of file phylo.hpp.

#### 7.19.3 Member Data Documentation

#### 7.19.3.1 blengths

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

Branch length.

Definition at line 29 of file phylo.hpp.

## 7.19.3.2 duplication

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

Definition at line 39 of file phylo.hpp.

#### 7.19.3.3 states

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

State of the parent node.

Definition at line 34 of file phylo.hpp.

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

• include/barry/counters/phylo.hpp

# 7.20 PhyloRuleDynData Class Reference

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

#### **Public Member Functions**

- PhyloRuleDynData (const std::vector< double > \*counts\_, uint pos\_, uint lb\_, uint ub\_, bool duplication\_)
- ∼PhyloRuleDynData ()

## **Public Attributes**

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

## 7.20.1 Detailed Description

Definition at line 1189 of file phylo.hpp.

#### 7.20.2 Constructor & Destructor Documentation

## 7.20.2.1 PhyloRuleDynData()

Definition at line 1196 of file phylo.hpp.

## 7.20.2.2 ~PhyloRuleDynData()

```
{\tt PhyloRuleDynData::}{\sim} {\tt PhyloRuleDynData ( ) [inline]}
```

Definition at line 1205 of file phylo.hpp.

## 7.20.3 Member Data Documentation

#### 7.20.3.1 counts

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

Definition at line 1191 of file phylo.hpp.

## 7.20.3.2 duplication

```
bool PhyloRuleDynData::duplication
```

Definition at line 1195 of file phylo.hpp.

#### 7.20.3.3 lb

```
uint PhyloRuleDynData::lb
```

Definition at line 1193 of file phylo.hpp.

## 7.20.3.4 pos

```
uint PhyloRuleDynData::pos
```

Definition at line 1192 of file phylo.hpp.

## 7.20.3.5 ub

```
uint PhyloRuleDynData::ub
```

Definition at line 1194 of file phylo.hpp.

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

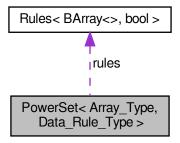
• include/barry/counters/phylo.hpp

#### 7.21 PowerSet < Array\_Type, Data\_Rule\_Type > Class Template Reference

Powerset of a binary array.

#include <powerset-bones.hpp>

Collaboration diagram for PowerSet < Array Type, Data Rule Type >:



## **Public Member Functions**

- void init\_support ()
- void calc ()
- void reset (uint N\_, uint M\_)

#### Construct and destroy a PowerSet object

- PowerSet ()
- PowerSet (uint N\_, uint M\_)
- PowerSet (const Array\_Type & array)
- ∼PowerSet ()

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

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

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

#### **Getter functions**

- const std::vector< Array\_Type > \* get\_data\_ptr () const
- std::vector< Array\_Type > get\_data () const
   std::vector< Array\_Type >::iterator begin ()
- std::vector< Array\_Type >::iterator end ()
- std::size\_t size () const noexcept
- const Array\_Type & operator[] (const unsigned int &i) const

#### **Public Attributes**

- Array\_Type EmptyArray
- std::vector< Array\_Type > data
- Rules< Array\_Type, Data\_Rule\_Type > \* rules
- uint N
- uint M
- bool rules\_deleted = false
- std::vector< std::pair< uint, uint >> coordinates\_free
- std::vector< std::pair< uint, uint >> coordinates\_locked

## 7.21.1 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.

#### 7.21.2 Constructor & Destructor Documentation

## 7.21.2.1 PowerSet() [1/3]

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

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

## 7.21.2.2 PowerSet() [2/3]

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

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

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

#### 7.21.2.4 ~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.

#### 7.21.3 Member Function Documentation

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

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

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

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

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

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

#### 7.21.3.4 begin()

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

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

#### 7.21.3.5 calc()

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

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

## 7.21.3.6 end()

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

Definition at line 74 of file powerset-bones.hpp.

## 7.21.3.7 get\_data()

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

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

#### 7.21.3.8 get\_data\_ptr()

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
const std::vector< Array_Type >* PowerSet< Array_Type, Data_Rule_Type >::get_data_ptr ()
const [inline]
```

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

#### 7.21.3.9 init\_support()

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

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

#### 7.21.3.10 operator[]()

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

## 7.21.3.11 reset()

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

#### 7.21.3.12 size()

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

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

## 7.21.4 Member Data Documentation

#### 7.21.4.1 coordinates free

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< std::pair<uint,uint> > PowerSet< Array_Type, Data_Rule_Type >::coordinates_free
```

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

#### 7.21.4.2 coordinates\_locked

template<typename Array\_Type = BArray<>, typename Data\_Rule\_Type = bool>
std::vector< std::pair<uint,uint> > PowerSet< Array\_Type, Data\_Rule\_Type >::coordinates\_←
locked

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

#### 7.21.4.3 data

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< Array_Type > PowerSet< Array_Type, Data_Rule_Type >::data
```

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

#### 7.21.4.4 EmptyArray

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

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

#### 7.21.4.5 M

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

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

## 7.21.4.6 N

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

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

#### 7.21.4.7 rules

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
Rules<Array_Type, Data_Rule_Type>* PowerSet< Array_Type, Data_Rule_Type >::rules
```

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

#### 7.21.4.8 rules\_deleted

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

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

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

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

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

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

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

## **Public Member Functions**

- ∼Rule ()
- Data\_Type \* D ()

Read/Write access to the data.

• bool operator() (const Array\_Type &a, uint i, uint j)

#### Construct a new Rule object

Construct a new Rule object

#### **Parameters**

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

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

## 7.22.1 Detailed Description

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

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

Rules can be used together with Support and PowerSet to determine which cells should be included when enumerating all possible realizations of a binary array.

#### **Template Parameters**

Array_Type	An object of class BArray.
Data_Type	Any type.

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

#### 7.22.2 Constructor & Destructor Documentation

#### 7.22.2.1 Rule() [1/2]

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

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

## 7.22.2.2 Rule() [2/2]

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

#### 7.22.2.3 ∼Rule()

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

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

#### 7.22.3 Member Function Documentation

#### 7.22.3.1 D()

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

Read/Write access to the data.

#### 7.22.3.2 operator()()

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

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

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

# 7.23 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 clear ()
- void get\_seq (const Array\_Type &a, std::vector< std::pair< uint, uint >> \*free, std::vector< std::pair< uint, uint >> \*locked=nullptr)

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

#### Rule adding

#### **Parameters**

rule

- void add rule (Rule < Array Type, Data Type > &rule)
- void add rule (Rule < Array Type, Data Type > \*rule)
- void add\_rule (Rule\_fun\_type< Array\_Type, Data\_Type > rule\_, Data\_Type \*data\_=nullptr, bool delete
   data =false)

# 7.23.1 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 69 of file rules-bones.hpp.

### 7.23.2 Constructor & Destructor Documentation

#### 7.23.2.1 Rules() [1/2]

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

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

# 7.23.2.2 Rules() [2/2]

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

#### 7.23.2.3 ∼Rules()

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

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

#### 7.23.3 Member Function Documentation

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

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

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

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

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

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

#### 7.23.3.4 clear()

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

Definition at line 127 of file rules-meat.hpp.

#### 7.23.3.5 get\_seq()

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

#### **Parameters**

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

#### Returns

Nothing.

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

#### 7.23.3.6 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 111 of file rules-meat.hpp.

# 7.23.3.7 operator=()

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

#### 7.23.3.8 size()

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

Definition at line 86 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

# 7.24 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 ()

Can be created without setting the array.

- ∼StatsCounter ()
- void reset\_array (const Array\_Type \*Array\_)

Changes the reference array for the counting.

- void add\_counter (Counter < Array\_Type, Data\_Type > \*f\_)
- void add\_counter (Counter< Array\_Type, Data\_Type > f\_)
- void set\_counters (Counters < Array\_Type, Data\_Type > \*counters\_)
- void count\_init (uint i, uint j)

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

- void count\_current (uint i, uint j)
- std::vector< double > count\_all ()
- Counters < Array\_Type, Data\_Type > \* get\_counters ()

# 7.24.1 Detailed Description

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

Count stats for a single Array.

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

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

#### 7.24.2 Constructor & Destructor Documentation

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

Creator of a StatsCounter

#### **Parameters**

Array⇔	A const pointer to a BArray.	
_		

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

# 7.24.2.2 StatsCounter() [2/2]

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

Can be created without setting the array.

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

# 7.24.2.3 ~StatsCounter()

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

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

#### 7.24.3 Member Function Documentation

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

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

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

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

#### 7.24.3.3 count all()

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

Definition at line 99 of file statscounter-meat.hpp.

#### 7.24.3.4 count\_current()

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

#### 7.24.3.5 count init()

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

Definition at line 61 of file statscounter-meat.hpp.

#### 7.24.3.6 get\_counters()

```
template<typename Array_Type , typename Data_Type >
Counters< Array_Type, Data_Type > * StatsCounter< Array_Type, Data_Type >::get_counters [inline]
```

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

#### 7.24.3.7 reset\_array()

Changes the reference array for the counting.

#### **Parameters**

Array⇔	A pointer to an array of class Array_Type.

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

#### 7.24.3.8 set\_counters()

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

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

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

# 7.25 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< std::vector< double > >
   \*stats\_bank=nullptr)
- void calc (std::vector < Array\_Type > \*array\_bank=nullptr, std::vector < std::vector < double > > \*stats\_←
  bank=nullptr, unsigned int max\_num\_elements\_=0u)

Computes the entire support.

- Counts\_type get\_counts () const
- const MapVec\_type \* get\_counts\_ptr () 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.
_	

- void reset\_array ()
- void reset\_array (const Array\_Type &Array\_)

#### **Manage counters**

#### **Parameters**

f_	A counter to be added.
counters←	A vector of counters to be added.
_	

- void add\_counter (Counter< Array\_Type, Data\_Counter\_Type > \*f\_)
- void add\_counter (Counter< Array\_Type, Data\_Counter\_Type > f\_)
- void set\_counters (Counters < Array\_Type, Data\_Counter\_Type > \*counters\_)

#### Manage rules

#### **Parameters**

f_	A rule to be added.
counters←	A vector of rules to be added.

- void add\_rule (Rule < Array\_Type, Data\_Rule\_Type > \*f\_)
- void add\_rule (Rule < Array\_Type, Data\_Rule\_Type > f\_)
- void set rules (Rules < Array Type, Data Rule Type > \*rules )
- 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\_)

### **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< std::pair< uint, uint >> coordinates free
- std::vector< std::pair< uint, uint >> coordinates\_locked
- std::vector< std::vector< double > > change\_stats

### 7.25.1 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\_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.

#### 7.25.2 Constructor & Destructor Documentation

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

Constructor passing a reference Array.

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

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

Constructor specifying the dimensions of the array (empty).

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

#### 7.25.2.3 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 85 of file support-bones.hpp.

#### 7.25.2.4 ~Support()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
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 92 of file support-bones.hpp.

#### 7.25.3 Member Function Documentation

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

Definition at line 204 of file support-meat.hpp.

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

Definition at line 214 of file support-meat.hpp.

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

Definition at line 241 of file support-meat.hpp.

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

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

#### 7.25.3.5 add\_rule\_dyn() [1/2]

Definition at line 276 of file support-meat.hpp.

### 7.25.3.6 add\_rule\_dyn() [2/2]

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

### 7.25.3.7 calc()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::calc (
    std::vector< Array_Type > * array_bank = nullptr,
    std::vector< std::vector< double > > * stats_bank = nullptr,
    unsigned int max_num_elements_ = 0u ) [inline]
```

Computes the entire support.

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

#### **Parameters**

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

Definition at line 178 of file support-meat.hpp.

# 7.25.3.8 get\_counters()

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

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

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

Vector of couter functions.

Definition at line 349 of file support-meat.hpp.

#### 7.25.3.9 get\_counts()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
Counts_type Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >
::get_counts [inline]
```

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

### 7.25.3.10 get\_counts\_ptr()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
const MapVec_type * Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type
>::get_counts_ptr [inline]
```

Definition at line 320 of file support-meat.hpp.

#### 7.25.3.11 get\_current\_stats()

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

List current statistics.

Definition at line 327 of file support-meat.hpp.

#### 7.25.3.12 get\_data()

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

Definition at line 344 of file support-meat.hpp.

#### 7.25.3.13 get\_rules()

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

Vector of static rules (cells to iterate).

Definition at line 354 of file support-meat.hpp.

#### 7.25.3.14 get\_rules\_dyn()

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

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

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

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

Definition at line 359 of file support-meat.hpp.

### 7.25.3.15 init\_support()

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

#### 7.25.3.16 print()

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

Definition at line 332 of file support-meat.hpp.

# 7.25.3.17 reset\_array() [1/2]

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

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

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

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

# 7.25.3.19 set\_counters()

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

# 7.25.3.20 set\_rules()

Definition at line 261 of file support-meat.hpp.

#### 7.25.3.21 set\_rules\_dyn()

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

void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::set_rules←
_dyn (

Rules< Array_Type, Data_Rule_Dyn_Type > * rules_ ) [inline]
```

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

#### 7.25.4 Member Data Documentation

#### 7.25.4.1 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< std::vector< double > > Support< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::change_stats
```

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

#### 7.25.4.2 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< std::pair<uint,uint> > Support< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::coordinates_free
```

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

#### 7.25.4.3 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< std::pair<uint,uint> > Support< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::coordinates_locked
```

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

#### 7.25.4.4 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 62 of file support-bones.hpp.

#### 7.25.4.5 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 56 of file support-bones.hpp.

#### 7.25.4.6 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 57 of file support-bones.hpp.

# 7.25.4.7 delete\_rules\_dyn

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_←
rules_dyn = true
```

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

#### 7.25.4.8 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 55 of file support-bones.hpp.

#### 7.25.4.9 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 59 of file support-bones.hpp.

#### 7.25.4.10 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 55 of file support-bones.hpp.

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

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

# 7.26 vecHasher < T > Struct Template Reference

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

#### **Public Member Functions**

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

# 7.26.1 Detailed Description

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

Definition at line 86 of file typedefs.hpp.

#### 7.26.2 Member Function Documentation

#### 7.26.2.1 operator()()

Definition at line 87 of file typedefs.hpp.

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

• include/barry/typedefs.hpp

# **Chapter 8**

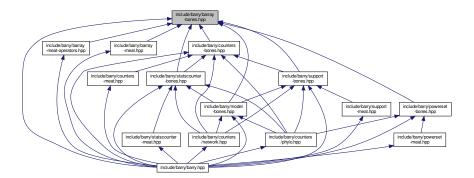
# **File Documentation**

# 8.1 include/barry/barray-bones.hpp File Reference

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



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



# Classes

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

#### **Macros**

• #define BARRAY\_BONES\_HPP 1

# 8.1.1 Macro Definition Documentation

# 8.1.1.1 BARRAY\_BONES\_HPP

#define BARRAY\_BONES\_HPP 1

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

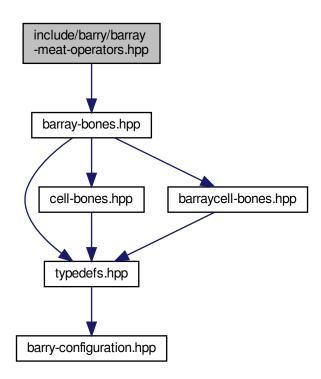
# 8.2 include/barry/barray-iterator.hpp File Reference

#### **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 ROW(a) this->el\_ij[a]
- #define COL(a) this->el\_ji[a]

#### **Functions**

template<typename Cell\_Type , typename Data\_Type >
 void checkdim\_ (const BArray< Cell\_Type, Data\_Type > &lhs, const BArray< Cell\_Type, Data\_Type > &rhs)

#### 8.3.1 Macro Definition Documentation

#### 8.3.1.1 BARRY\_BARRAY\_MEAT\_OPERATORS\_HPP

```
#define BARRY_BARRAY_MEAT_OPERATORS_HPP 1
```

Definition at line 5 of file barray-meat-operators.hpp.

#### 8.3.1.2 COL

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

#### 8.3.1.3 ROW

Definition at line 7 of file barray-meat-operators.hpp.

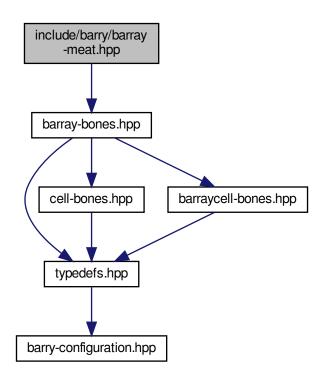
#### 8.3.2 Function Documentation

# 8.3.2.1 checkdim\_()

Definition at line 11 of file barray-meat-operators.hpp.

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

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



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



# **Macros**

- #define ROW(a) this->el\_ij[a]
- #define COL(a) this->el\_ji[a]

#### 8.4.1 Macro Definition Documentation

#### 8.4.1.1 COL

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

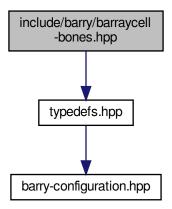
# 8.4.1.2 ROW

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

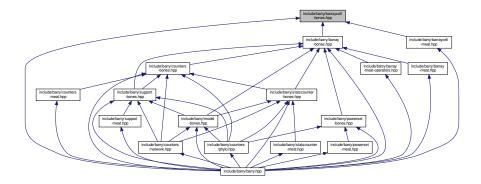
Definition at line 7 of file barray-meat.hpp.

# 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:

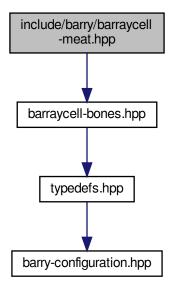


# Classes

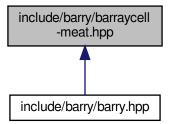
- class BArrayCell
   Cell\_Type, Data\_Type
- class BArrayCell\_const< Cell\_Type, Data\_Type >

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

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

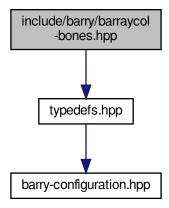


This graph shows which files directly or indirectly include this file:

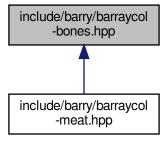


# 8.7 include/barry/barraycol-bones.hpp File Reference

#include "typedefs.hpp"
Include dependency graph for barraycol-bones.hpp:



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

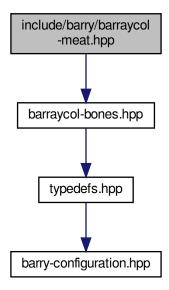


#### **Classes**

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

# 8.8 include/barry/barraycol-meat.hpp File Reference

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



#### **Macros**

• #define BARRY\_BARRAYCOL\_MEAT\_HPP 1

#### 8.8.1 Macro Definition Documentation

# 8.8.1.1 BARRY\_BARRAYCOL\_MEAT\_HPP

#define BARRY\_BARRAYCOL\_MEAT\_HPP 1

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

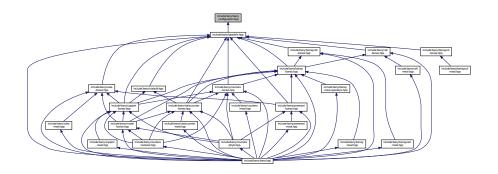
# 8.9 include/barry/barraydense-bones.hpp File Reference

#### **Classes**

class BArrayDense < Cell\_Type, Data\_Type >
 Dense bi-dimensional array.

# 8.10 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.

```
• #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< unsigned int >(UINT\_MAX/2u)
- template<typename Ta , typename Tb >
   using Map = std::map< Ta, Tb >

#### 8.10.1 Macro Definition Documentation

#### 8.10.1.1 BARRY\_CHECK\_SUPPORT

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

Definition at line 45 of file barry-configuration.hpp.

# 8.10.1.2 BARRY\_ISFINITE

Definition at line 38 of file barry-configuration.hpp.

#### 8.10.1.3 BARRY\_MAX\_NUM\_ELEMENTS

```
\#define BARRY_MAX_NUM_ELEMENTS static_cast< unsigned int >(UINT_MAX/2u)
```

Definition at line 53 of file barry-configuration.hpp.

# 8.10.1.4 BARRY\_SAFE\_EXP

```
#define BARRY_SAFE_EXP -100.0
```

Definition at line 31 of file barry-configuration.hpp.

#### 8.10.1.5 printf\_barry

```
#define printf_barry printf
```

Definition at line 49 of file barry-configuration.hpp.

# 8.10.2 Typedef Documentation

#### 8.10.2.1 Map

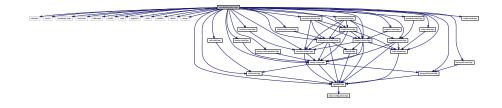
```
template<typename Ta , typename Tb >
using Map = std::map<Ta,Tb>
```

Definition at line 25 of file barry-configuration.hpp.

# 8.11 include/barry/barry.hpp File Reference

```
#include <iostream>
#include <vector>
#include <unordered_map>
#include <functional>
#include <stdexcept>
#include <cmath>
#include <map>
#include <algorithm>
#include <utility>
#include <random>
#include <climits>
#include <string>
#include "typedefs.hpp"
#include "cell-bones.hpp"
#include "cell-meat.hpp"
#include "barray-bones.hpp"
#include "barraycell-bones.hpp"
#include "barray-meat.hpp"
#include "barraycell-meat.hpp"
#include "barray-meat-operators.hpp"
#include "counters-bones.hpp"
#include "counters-meat.hpp"
#include "statscounter-bones.hpp"
```

```
#include "statscounter-meat.hpp"
#include "support-bones.hpp"
#include "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)

#### 8.11.1 Macro Definition Documentation

#### 8.11.1.1 BARRY HPP

#define BARRY\_HPP

Definition at line 19 of file barry.hpp.

# 8.11.1.2 BARRY\_VERSION

```
#define BARRY_VERSION 0.1
```

Definition at line 21 of file barry.hpp.

#### 8.11.1.3 COUNTER\_FUNCTION

```
\#define COUNTER_FUNCTION( a )
```

#### Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
inline double (a) (const Array_Type & Array, uint i, uint j, Data_Type * data) \
```

Definition at line 72 of file barry.hpp.

#### 8.11.1.4 COUNTER LAMBDA

```
#define 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 75 of file barry.hpp.

# 8.11.1.5 RULE\_FUNCTION

```
#define RULE_FUNCTION( a )
```

#### Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
inline bool (a) (const Array_Type & Array, uint i, uint j, Data_Type * data)\
```

Definition at line 79 of file barry.hpp.

#### 8.11.1.6 RULE\_LAMBDA

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

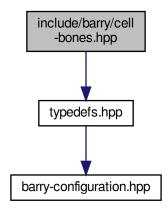
#### Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
Rule_fun_type<Array_Type, Data_Type> a = \
[](const Array_Type & Array, uint i, uint j, Data_Type * data)
```

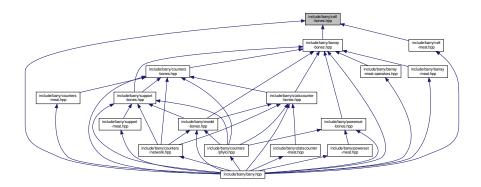
Definition at line 82 of file barry.hpp.

# 8.12 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:



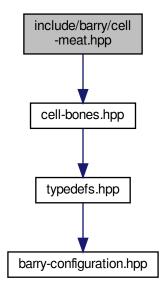
# Classes

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

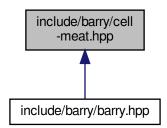
# 8.13 include/barry/cell-meat.hpp File Reference

#include "cell-bones.hpp"

Include dependency graph for cell-meat.hpp:



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

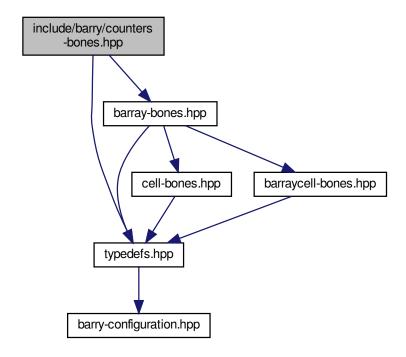


# 8.14 include/barry/col-bones.hpp File Reference

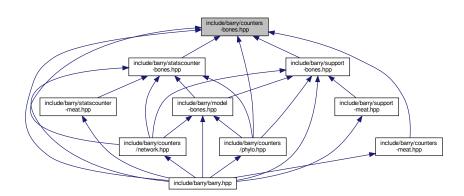
# 8.15 include/barry/counters-bones.hpp File Reference

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

Include dependency graph for counters-bones.hpp:



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

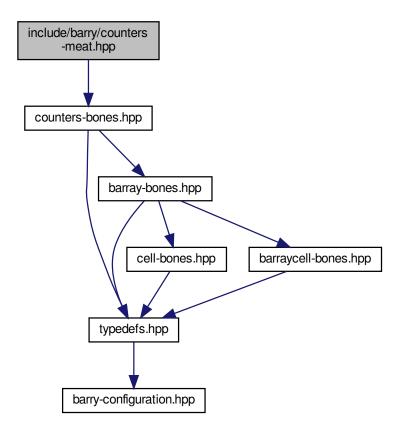


# **Classes**

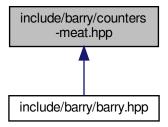
- class Counter < Array\_Type, Data\_Type >
   A counter function based on change statistics.

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

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

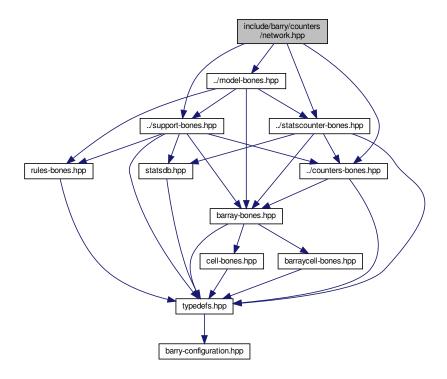


This graph shows which files directly or indirectly include this file:

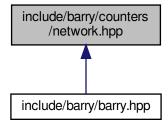


# 8.17 include/barry/counters/network.hpp File Reference

```
#include "../counters-bones.hpp"
#include "../support-bones.hpp"
#include "../statscounter-bones.hpp"
#include "../model-bones.hpp"
Include dependency graph for network.hpp:
```



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



# **Classes**

class NetworkData

Data class for Networks.

class NetCounterData

Data class used to store arbitrary uint or double vectors.

#### **Macros**

- #define NET\_C\_DATA\_IDX(i) (data->indices[i])
- #define NET\_C\_DATA\_NUM(i) (data->numbers[i])

#### Macros for defining counters

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

#### Macros for defining rules

- #define NETWORK\_RULE(a)
- #define NETWORK\_RULE\_LAMBDA(a)

# **Typedefs**

#### Convenient typedefs for network objects.

- typedef BArray< double, NetworkData > Network
- typedef Counter
   NetWork, NetCounterData > NetCounter
- typedef Counters < Network, NetCounterData > NetCounters
- typedef Support < Network, NetCounterData > NetSupport
- typedef StatsCounter
   NetWork, NetCounterData > NetStatsCounter
- typedef Model < Network, NetCounterData > NetModel
- typedef Rule < Network, bool > NetRule
- typedef Rules < Network, bool > NetRules

## **Functions**

void counter edges (NetCounters \*counters)

Number of edges.

• void counter\_isolates (NetCounters \*counters)

Number of isolated vertices.

void counter\_mutual (NetCounters \*counters)

Number of mutual ties.

- void counter\_istar2 (NetCounters \*counters)
- void counter ostar2 (NetCounters \*counters)
- void counter ttriads (NetCounters \*counters)
- void counter\_ctriads (NetCounters \*counters)
- void counter\_density (NetCounters \*counters)
- void counter\_idegree15 (NetCounters \*counters)
- void counter\_odegree15 (NetCounters \*counters)
- void counter\_absdiff (NetCounters \*counters, uint attr\_id, double alpha=1.0)

Sum of absolute attribute difference between ego and alter.

- void counter\_diff (NetCounters \*counters, uint attr\_id, double alpha=1.0, double tail\_head=true)

  Sum of attribute difference between ego and alter to pow(alpha)
- NETWORK COUNTER (init single attr)
- void counter\_nodeicov (NetCounters \*counters, uint attr\_id)
- void counter\_nodeocov (NetCounters \*counters, uint attr\_id)
- void counter nodecov (NetCounters \*counters, uint attr id)
- void counter\_nodematch (NetCounters \*counters, uint attr\_id)
- void counter\_idegree (NetCounters \*counters, std::vector< uint > d)

Counts number of vertices with a given in-degree.

void counter\_odegree (NetCounters \*counters, std::vector< uint > d)

Counts number of vertices with a given out-degree.

void counter\_degree (NetCounters \*counters, std::vector< uint > d)

Counts number of vertices with a given out-degree.

#### Rules for network models

#### **Parameters**

```
rules | A pointer to a NetRules object (Rules < Network, bool > ).
```

void rules\_zerodiag (NetRules \*rules)
 Number of edges.

## 8.17.1 Macro Definition Documentation

# 8.17.1.1 NET\_C\_DATA\_IDX

Definition at line 79 of file network.hpp.

# 8.17.1.2 NET C DATA NUM

Definition at line 80 of file network.hpp.

# 8.17.1.3 NETWORK\_COUNTER

Function for definition of a network counter function

Definition at line 101 of file network.hpp.

## 8.17.1.4 NETWORK\_COUNTER\_LAMBDA

Lambda function for definition of a network counter function

Definition at line 104 of file network.hpp.

## 8.17.1.5 NETWORK\_RULE

Function for definition of a network counter function

Definition at line 113 of file network.hpp.

## 8.17.1.6 NETWORK\_RULE\_LAMBDA

Lambda function for definition of a network counter function

Definition at line 116 of file network.hpp.

# 8.17.2 Typedef Documentation

#### 8.17.2.1 NetCounter

typedef Counter<Network, NetCounterData > NetCounter

Definition at line 88 of file network.hpp.

#### 8.17.2.2 NetCounters

typedef Counters< Network, NetCounterData> NetCounters

Definition at line 89 of file network.hpp.

#### 8.17.2.3 NetModel

typedef Model < Network, NetCounterData > NetModel

Definition at line 92 of file network.hpp.

#### 8.17.2.4 NetRule

typedef Rule<Network,bool> NetRule

Definition at line 93 of file network.hpp.

#### 8.17.2.5 NetRules

typedef Rules<Network,bool> NetRules

Definition at line 94 of file network.hpp.

## 8.17.2.6 NetStatsCounter

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

Definition at line 91 of file network.hpp.

## 8.17.2.7 NetSupport

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

Definition at line 90 of file network.hpp.

#### 8.17.2.8 Network

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

Definition at line 87 of file network.hpp.

# 8.17.3 Function Documentation

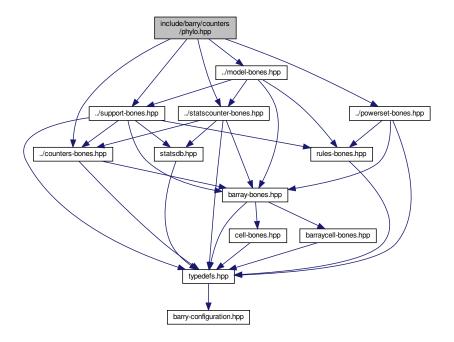
# 8.17.3.1 rules\_zerodiag()

Number of edges.

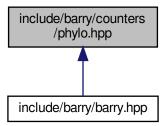
Definition at line 742 of file network.hpp.

# 8.18 include/barry/counters/phylo.hpp File Reference

```
#include "../counters-bones.hpp"
#include "../support-bones.hpp"
#include "../statscounter-bones.hpp"
#include "../model-bones.hpp"
#include "../powerset-bones.hpp"
Include dependency graph for phylo.hpp:
```



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



## **Classes**

- class NodeData
   Data definition for the PhyloArray class.
- class PhyloRuleDynData

#### **Macros**

#define PHYLO\_COUNTER\_LAMBDA(a)

Extension of a simple counter.

- #define PHYLO RULE DYN LAMBDA(a)
- #define PHYLO\_CHECK\_MISSING()

## **Typedefs**

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

#### Convenient typedefs for Node objects.

- typedef BArray< uint, NodeData > PhyloArray
- typedef Counter< PhyloArray, PhyloCounterData > PhyloCounter
- typedef Counters
   PhyloArray, PhyloCounterData > PhyloCounters
- typedef Rule < PhyloArray, PhyloRuleData > PhyloRule
- typedef Rules< PhyloArray, PhyloRuleData > PhyloRules
- typedef 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 (bool d)
- void counter\_overall\_gains (PhyloCounters \*counters, bool duplication=true)

Overall functional gains.

void counter gains (PhyloCounters \*counters, std::vector< uint > nfun, bool duplication=true)

Functional gains for a specific function (nfun).

void counter\_gains\_k\_offspring (PhyloCounters \*counters, std::vector< uint > nfun, uint k=1u, bool duplication=true)

k genes gain function nfun

void counter\_genes\_changing (PhyloCounters \*counters, bool duplication=true)

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

• void counter\_overall\_loss (PhyloCounters \*counters, bool duplication=true)

Overall functional loss.

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

Cap the number of functions per gene.

void counter\_loss (PhyloCounters \*counters, std::vector< uint > nfun, bool duplication=true)

Total count of losses for an specific function.

• void counter\_overall\_changes (PhyloCounters \*counters, bool duplication=true)

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

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

Total count of Sub-functionalization events.

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

Co-evolution (joint gain or loss)

void counter\_longest (PhyloCounters \*counters)

Longest branch mutates (either by gain or by loss)

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

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

  Total number of neofunctionalization events.
- void counter\_co\_opt (PhyloCounters \*counters, uint nfunA, uint nfunB, bool duplication=true) Function co-opting.
- void rule\_dyn\_limit\_changes (PhyloSupport \*support, uint pos, uint lb, uint ub, bool duplication=true)

  Overall functional gains.

## 8.18.1 Macro Definition Documentation

## 8.18.1.1 PHYLO\_CHECK\_MISSING

```
#define PHYLO_CHECK_MISSING( )

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

Definition at line 94 of file phylo.hpp.

## 8.18.1.2 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 88 of file phylo.hpp.

# 8.18.1.3 PHYLO\_RULE\_DYN\_LAMBDA

## Value:

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

Definition at line 91 of file phylo.hpp.

# 8.18.2 Typedef Documentation

## 8.18.2.1 PhyloArray

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

Definition at line 61 of file phylo.hpp.

# 8.18.2.2 PhyloCounter

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

Definition at line 62 of file phylo.hpp.

# 8.18.2.3 PhyloCounterData

```
typedef std::vector< uint > PhyloCounterData
```

Definition at line 53 of file phylo.hpp.

# 8.18.2.4 PhyloCounters

```
typedef Counters< PhyloArray, PhyloCounterData> PhyloCounters
```

Definition at line 63 of file phylo.hpp.

# 8.18.2.5 PhyloModel

typedef Model<PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloModel

Definition at line 73 of file phylo.hpp.

## 8.18.2.6 PhyloPowerSet

typedef PowerSet<PhyloArray, PhyloRuleData> PhyloPowerSet

Definition at line 74 of file phylo.hpp.

## 8.18.2.7 PhyloRule

typedef Rule<PhyloArray,PhyloRuleData> PhyloRule

Definition at line 65 of file phylo.hpp.

# 8.18.2.8 PhyloRuleData

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

Definition at line 54 of file phylo.hpp.

# 8.18.2.9 PhyloRuleDyn

typedef Rule<PhyloArray,PhyloRuleDynData> PhyloRuleDyn

Definition at line 68 of file phylo.hpp.

## 8.18.2.10 PhyloRules

typedef Rules<PhyloArray,PhyloRuleData> PhyloRules

Definition at line 66 of file phylo.hpp.

# 8.18.2.11 PhyloRulesDyn

```
typedef Rules<PhyloArray,PhyloRuleDynData> PhyloRulesDyn
```

Definition at line 69 of file phylo.hpp.

# 8.18.2.12 PhyloStatsCounter

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

Definition at line 72 of file phylo.hpp.

## 8.18.2.13 PhyloSupport

```
{\tt typedef~Support < PhyloArray,~PhyloCounterData,~PhyloRuleData,~PhyloRuleDynData > PhyloSupport}
```

Definition at line 71 of file phylo.hpp.

## 8.18.3 Function Documentation

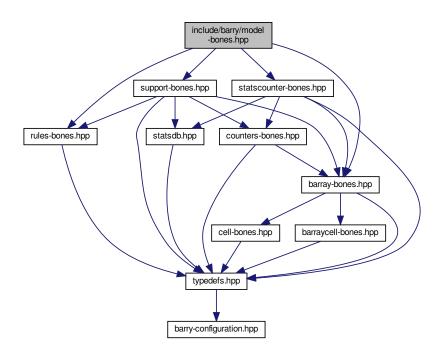
# 8.18.3.1 get\_last\_name()

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

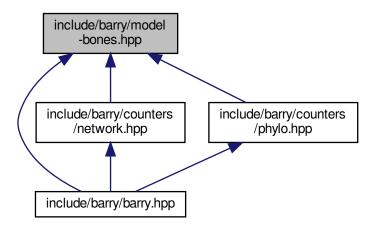
Definition at line 99 of file phylo.hpp.

# 8.19 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**

- double update\_normalizing\_constant (const std::vector< double > &params, const Counts\_type &support)
- double likelihood\_ (const std::vector< double > &target\_stats, const std::vector< double > &params, const double normalizing\_constant, bool log\_=false)
- template<typename Array\_Type >
   std::vector< double > keygen\_default (const Array\_Type &Array\_)

Array Hasher class (used for computing support)

#### 8.19.1 Function Documentation

## 8.19.1.1 keygen\_default()

Array Hasher class (used for computing support)

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

## 8.19.1.2 likelihood\_()

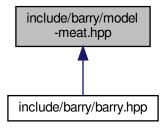
Definition at line 40 of file model-bones.hpp.

#### 8.19.1.3 update\_normalizing\_constant()

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

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

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



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

```
#include "geese/geese-node-bones.hpp"
#include "geese/geese-bones.hpp"
#include "geese/geese-meat.hpp"
#include "geese/geese-meat-constructors.hpp"
#include "geese/geese-meat-likelihood.hpp"
#include "geese/geese-meat-likelihood_exhaust.hpp"
#include "geese/geese-meat-simulate.hpp"
#include "geese/geese-meat-predict.hpp"
#include "geese/flock-bones.hpp"
#include "geese/flock-meet.hpp"
Include dependency graph for geese.hpp:
```



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

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



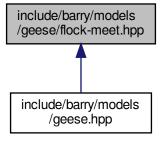
# Classes

· class Flock

A Flock is a group of Geese.

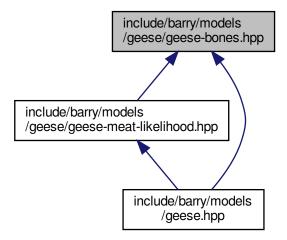
# 8.23 include/barry/models/geese/flock-meet.hpp File Reference

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



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

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



# **Classes**

• class Geese

Annotated Phylo Model.

#### **Macros**

• #define INITIALIZED()

# **Functions**

- template<typename Ta , typename Tb >  $std::vector < Ta > vector\_caster \ (const \ std::vector < Tb > \&x)$
- RULE\_FUNCTION (rule\_empty\_free)
- std::vector< double > keygen\_full (const phylocounters::PhyloArray &array)
- bool vec\_diff (const std::vector< unsigned int > &s, const std::vector< unsigned int > &a)

# 8.24.1 Macro Definition Documentation

#### 8.24.1.1 INITIALIZED

```
#define INITIALIZED( )

Value:
    if (!this->initialized) \
        throw std::logic_error("The model has not been initialized yet.");
```

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

# 8.24.2 Function Documentation

## 8.24.2.1 keygen\_full()

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

# 8.24.2.2 RULE\_FUNCTION()

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

## 8.24.2.3 vec\_diff()

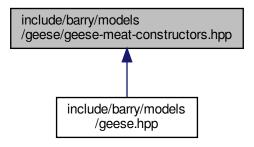
Definition at line 51 of file geese-bones.hpp.

# 8.24.2.4 vector\_caster()

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

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

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

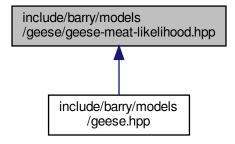


# 8.26 include/barry/models/geese/geese-meat-likelihood.hpp File Reference

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

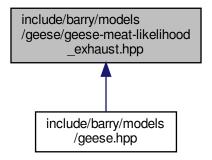


This graph shows which files directly or indirectly include this file:



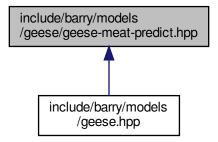
# 8.27 include/barry/models/geese/geese-meat-likelihood\_exhaust.hpp File Reference

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



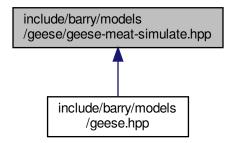
# 8.28 include/barry/models/geese/geese-meat-predict.hpp File Reference

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



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

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



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

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



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

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



## Classes

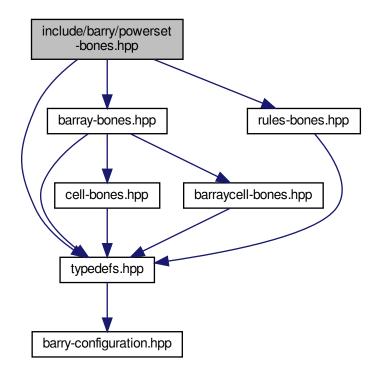
• class Node

A single node for the model.

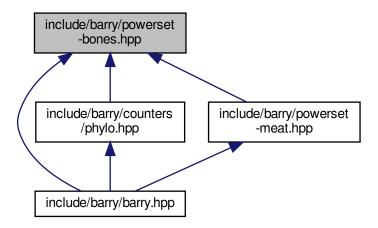
# 8.32 include/barry/powerset-bones.hpp File Reference

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

Include dependency graph for powerset-bones.hpp:



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

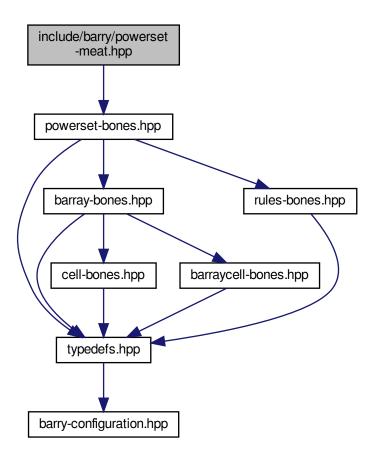


# **Classes**

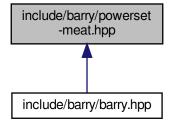
# 8.33 include/barry/powerset-meat.hpp File Reference

#include "powerset-bones.hpp"

Include dependency graph for powerset-meat.hpp:

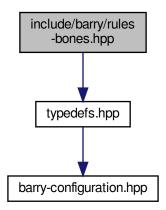


This graph shows which files directly or indirectly include this file:

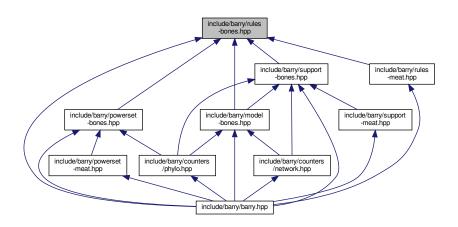


# 8.34 include/barry/rules-bones.hpp File Reference

#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)
```

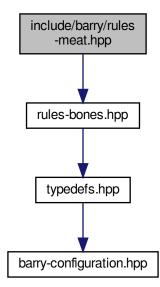
## 8.34.1 Function Documentation

## 8.34.1.1 rule\_fun\_default()

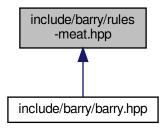
Definition at line 10 of file rules-bones.hpp.

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

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



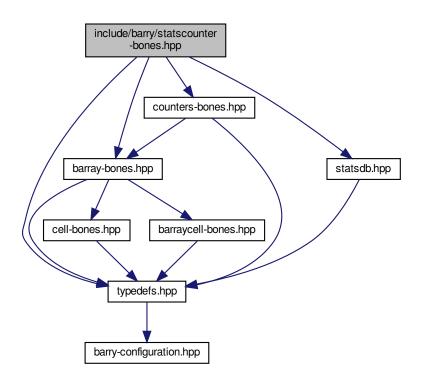
This graph shows which files directly or indirectly include this file:



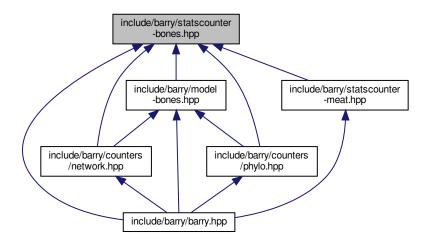
# 8.36 include/barry/statscounter-bones.hpp File Reference

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

Include dependency graph for statscounter-bones.hpp:



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



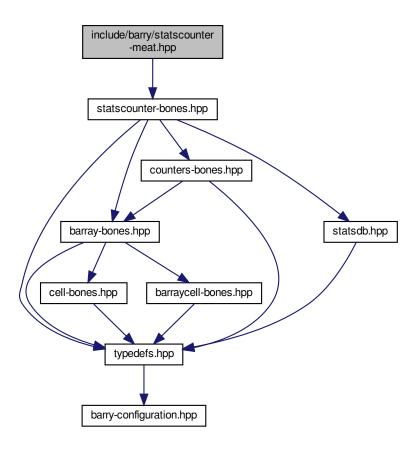
# Classes

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

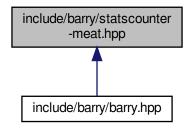
# 8.37 include/barry/statscounter-meat.hpp File Reference

#include "statscounter-bones.hpp"

Include dependency graph for statscounter-meat.hpp:



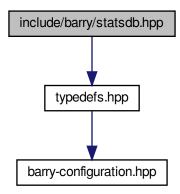
This graph shows which files directly or indirectly include this file:



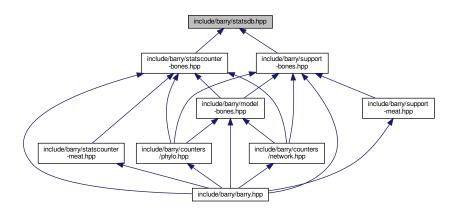
# 8.38 include/barry/statsdb.hpp File Reference

#include "typedefs.hpp"

Include dependency graph for statsdb.hpp:



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



## **Classes**

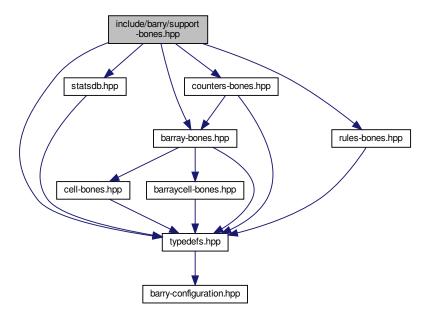
class FreqTable < T >

Database of statistics.

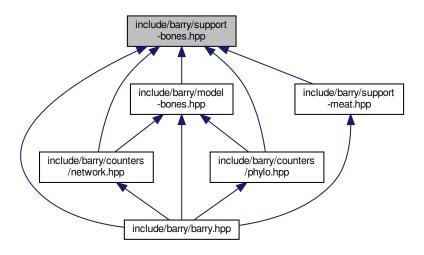
# 8.39 include/barry/support-bones.hpp File Reference

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

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



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

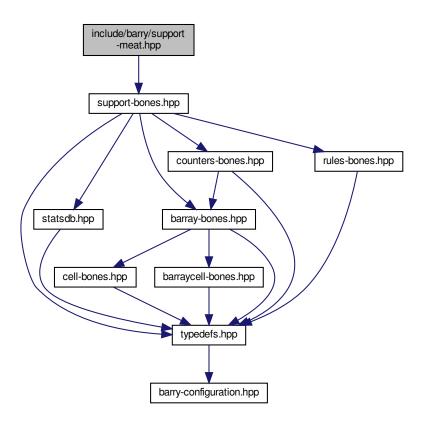


# **Classes**

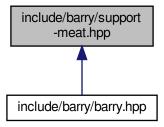
class Support < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type >
 Compute the support of sufficient statistics.

# 8.40 include/barry/support-meat.hpp File Reference

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



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



#### **Macros**

• #define BARRY\_SUPPORT\_MEAT\_HPP 1

## 8.40.1 Macro Definition Documentation

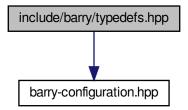
## 8.40.1.1 BARRY SUPPORT MEAT HPP

#define BARRY\_SUPPORT\_MEAT\_HPP 1

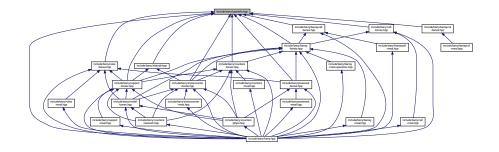
Definition at line 4 of file support-meat.hpp.

# 8.41 include/barry/typedefs.hpp File Reference

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



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



#### **Classes**

class Entries < Cell\_Type >

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

struct vecHasher< T >

## **Namespaces**

CHECK

Integer constants used to specify which cell should be check.

• EXISTS

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

# **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 >
        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-10)
```

## **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

# 8.41.1 Typedef Documentation

#### 8.41.1.1 Col type

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

Definition at line 51 of file typedefs.hpp.

## 8.41.1.2 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 123 of file typedefs.hpp.

#### 8.41.1.3 Counts\_type

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

Definition at line 44 of file typedefs.hpp.

## 8.41.1.4 MapVec\_type

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

Definition at line 105 of file typedefs.hpp.

## 8.41.1.5 Row\_type

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

Definition at line 48 of file typedefs.hpp.

#### 8.41.1.6 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 126 of file typedefs.hpp.

## 8.41.1.7 uint

```
typedef unsigned int uint
```

Definition at line 10 of file typedefs.hpp.

## 8.41.2 Function Documentation

# 8.41.2.1 vec\_equal()

Compares if -a- and -b- are equal.

## **Parameters**

```
a,b Two vectors of the same length
```

#### Returns

true if all elements are equal.

Definition at line 137 of file typedefs.hpp.

198 File Documentation

# 8.41.2.2 vec\_equal\_approx()

Definition at line 155 of file typedefs.hpp.

# 8.41.2.3 vec\_inner\_prod()

Definition at line 175 of file typedefs.hpp.

# 8.42 README.md File Reference

# Index

```
\simBArray
                                                          StatsCounter< Array Type, Data Type >, 129
                                                     \sim\!\!\text{Support}
    BArray< Cell_Type, Data_Type >, 31
                                                                       Array_Type,
                                                                                      Data Counter Type,
\simBArrayCell
                                                          Support<
    BArrayCell< Cell_Type, Data_Type >, 42
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
~BArrayCell const
                                                               135
    BArrayCell_const< Cell_Type, Data_Type >, 44
                                                     add
\simBArrayCol
                                                          Cell< Cell_Type >, 57
    BArrayCol < Cell_Type, Data_Type >, 47
                                                          FreqTable < T >, 79
~BArrayCol const
                                                     add_array
    BArrayCol_const< Cell_Type, Data_Type >, 50
                                                          Model <
                                                                      Array_Type,
                                                                                      Data_Counter_Type,
\simBArrayDense
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
    BArrayDense < Cell Type, Data Type >, 52
                                                               93
\simCell
                                                      add counter
    Cell< Cell_Type >, 56
                                                          Counters < Array_Type, Data_Type >, 68
\simConstBArrayRowIter
                                                          Model<
                                                                      Array_Type,
                                                                                      Data Counter Type,
    ConstBArrayRowlter< Cell Type, Data Type >, 60
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
\simCounter
                                                               94
    Counter< Array_Type, Data_Type >, 63
                                                          StatsCounter < Array_Type, Data_Type >, 129
\simCounters
                                                                                      Data_Counter_Type,
                                                          Support<
                                                                       Array_Type,
    Counters < Array_Type, Data_Type >, 67
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
\simEntries
    Entries < Cell Type >, 71
                                                      add data
\simFlock
                                                          Flock, 74
    Flock, 74
                                                     add rule
\simFreqTable
                                                          Model<
                                                                      Array Type,
                                                                                      Data Counter Type,
    FreqTable < T >, 79
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
\simGeese
                                                               94.95
    Geese, 84
                                                          PowerSet < Array_Type, Data_Rule_Type >, 118
\simModel
                                                          Rules < Array_Type, Data_Type >, 126
    Model<
                Array_Type,
                                Data_Counter_Type,
                                                          Support<
                                                                      Array_Type,
                                                                                      Data Counter Type,
         Data_Rule_Type, Data_Rule_Dyn_Type >,
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
         93
                                                               135, 136
\simNetCounterData
                                                      add rule dyn
    NetCounterData, 103
                                                          Model<
                                                                      Array_Type,
                                                                                      Data Counter Type,
\simNetworkData
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
    NetworkData, 105
                                                               95, 96
\simNode
                                                          Support<
                                                                       Array_Type,
                                                                                      Data Counter Type,
    Node, 108
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
\simNodeData
    NodeData, 112
                                                     annotations
\simPhyloRuleDynData
                                                          Node, 109
    PhyloRuleDynData, 114
                                                     Array
\simPowerSet
                                                          ConstBArrayRowlter< Cell Type, Data Type >, 60
    PowerSet< Array_Type, Data_Rule_Type >, 118
                                                     array
\simRule
                                                          Node, 109
    Rule < Array_Type, Data_Type >, 123
                                                     arrays
\simRules
                                                          Node, 109
    Rules < Array_Type, Data_Type >, 125
                                                      AS ONE
\simStatsCounter
                                                          EXISTS, 25
```

as vector	COL 140
as_vector FreqTable < T >, 79	COL, 148 ROW, 148
AS ZERO	BARRAY_BONES_HPP
EXISTS, 25	barray-bones.hpp, 144
LXI010, 20	BArrayCell
BArray	BArrayCell< Cell Type, Data Type >, 42
BArray< Cell_Type, Data_Type >, 30, 31	BArrayCell< Cell_Type, Data_Type >, 41
BArray< Cell_Type, Data_Type >, 27	~BArrayCell, 42
∼BArray, <mark>31</mark>	BArray Cell Type, Data Type >, 40
BArray, 30, 31	BArrayCell, 42
BArrayCell< Cell_Type, Data_Type >, 40	operator Cell_Type, 42
BArrayCell_const< Cell_Type, Data_Type >, 40	operator*=, 42
clear, 31	operator+=, 42
col, 31	operator-=, 43
D, 32	operator/=, 43
default_val, 32	operator=, 43
get_cell, 32	operator==, 43
get_col, 32	BArrayCell_const
get_col_vec, 32, 33	BArrayCell const< Cell Type, Data Type >, 44
get_entries, 33	BArrayCell const< Cell Type, Data Type >, 44
get_row, 33	~BArrayCell_const, 44
get_row_vec, 33	BArray < Cell_Type, Data_Type >, 40
insert_cell, 34	BArrayCell_const, 44
is_empty, 34	operator Cell_Type, 45
ncol, 34	operator!=, 45
nnozero, 35	operator<, 45
nrow, 35	operator<=, 45
operator*=, 35	operator>, 45
operator(), 35	operator>=, 46
operator+=, 35, 36	operator==, 45
operator-=, 36	BArrayCol
operator/=, 36	BArrayCol< Cell_Type, Data_Type >, 47
operator=, 37	BArrayCol_const< Cell_Type, Data_Type >, 50
operator==, 37 out of range, 37	BArrayCol < Cell_Type, Data_Type >, 46
print, 37	~BArrayCol, 47
reserve, 37	BArrayCol, 47
resize, 38	begin, 47
rm_cell, 38	end, 47
row, 38	operator Cell_Type, 47
set_data, 38	operator*=, 48
swap_cells, 39	operator+=, 48
swap cols, 39	operator-=, 48 operator/=, 48
swap_rows, 39	operator=, 48
toggle_cell, 39	operator==, 49
toggle_lock, 39	barraycol-meat.hpp
transpose, 40	BARRY_BARRAYCOL_MEAT_HPP, 152
visited, 41	BArrayCol_const< Cell_Type, Data_Type >, 49
zero_col, 40	~BArrayCol_const, 50
zero_row, 40	BArrayCol, 50
barray-bones.hpp	operator!=, 50
BARRAY_BONES_HPP, 144	operator<, 50
barray-meat-operators.hpp	operator<=, 50
BARRY_BARRAY_MEAT_OPERATORS_HPP,	operator>, 51
146	operator>=, 51
checkdim_, 146	operator==, 51
COL, 146	BArrayDense
ROW, 146	BArrayDense < Cell_Type, Data_Type >, 52
barray-meat.hpp	

BArrayDense< Cell_Type, Data_Type >, 51	Support< Array_Type, Data_Counter_Type,
$\sim$ BArrayDense, 52	Data_Rule_Type, Data_Rule_Dyn_Type >,
BArrayDense, 52	136
elements_ptr, 53	calc_reduced_sequence
elements_raw, 53	Geese, 84
fill, 53	calc_sequence
ncol, 53	Geese, 84
nrow, 53	Cell
operator(), 54	Cell< Cell Type >, 55-57
operator(), 54	Cell< Cell_Type >, 54
	— ··
print, 54	~Cell, 56
barry, 23	add, 57
barry-configuration.hpp	Cell, 55–57
BARRY_CHECK_SUPPORT, 153	operator Cell_Type, 58
BARRY_ISFINITE, 153	operator=, 58
BARRY_MAX_NUM_ELEMENTS, 153	value, 58
BARRY_SAFE_EXP, 153	visited, 58
Map, 154	change_stats
printf_barry, 154	Support< Array_Type, Data_Counter_Type,
barry.hpp	Data_Rule_Type, Data_Rule_Dyn_Type >,
BARRY HPP, 155	140
BARRY_VERSION, 155	CHECK, 24
COUNTER_FUNCTION, 156	BOTH, 24
COUNTER_LAMBDA, 156	NONE, 24
RULE_FUNCTION, 156	ONE, 24
RULE_LAMBDA, 156	TWO, 24
barry::counters, 23	checkdim_
barry::counters::network, 24	barray-meat-operators.hpp, 146
barry::counters::phylo, 24	clear
BARRY_BARRAY_MEAT_OPERATORS_HPP	BArray< Cell_Type, Data_Type >, 31
barray-meat-operators.hpp, 146	Counters< Array_Type, Data_Type >, 68
BARRY_BARRAYCOL_MEAT_HPP	FreqTable $< T >$ , 80
	•
barraycol-meat.hpp, 152	Rules < Array_Type, Data_Type >, 126
BARRY_CHECK_SUPPORT	COL
barry-configuration.hpp, 153	barray-meat-operators.hpp, 146
BARRY_HPP	barray-meat.hpp, 148
barry.hpp, 155	col
BARRY_ISFINITE	BArray< Cell_Type, Data_Type >, 31
barry-configuration.hpp, 153	Col_type
BARRY_MAX_NUM_ELEMENTS	typedefs.hpp, 196
barry-configuration.hpp, 153	ConstBArrayRowlter
BARRY_SAFE_EXP	ConstBArrayRowlter< Cell_Type, Data_Type >, 60
barry-configuration.hpp, 153	ConstBArrayRowlter< Cell_Type, Data_Type >, 59
BARRY_SUPPORT_MEAT_HPP	~ConstBArrayRowlter, 60
	-
support-meat.hpp, 194	Array, 60
BARRY_VERSION	ConstBArrayRowlter, 60
barry.hpp, 155	current_col, 60
begin	current_row, 60
BArrayCol < Cell_Type, Data_Type >, 47	iter, 61
PowerSet< Array_Type, Data_Rule_Type >, 118	coordinates_free
blengths	PowerSet< Array_Type, Data_Rule_Type >, 120
NodeData, 113	Support< Array_Type, Data_Counter_Type,
ВОТН	Data_Rule_Type, Data_Rule_Dyn_Type >,
CHECK, 24	140
EXISTS, 25	coordinates_locked
calo	PowerSet < Array_Type, Data_Rule_Type >, 120
Calc	Support< Array_Type, Data_Counter_Type,
PowerSet < Array_Type, Data_Rule_Type >, 119	Data Rule Type. Data Rule Dyn Type >.

140	Network counters, 12
count	COUNTER_LAMBDA
Counter< Array_Type, Data_Type >, 63	barry.hpp, 156
count_all	counter_longest
StatsCounter< Array_Type, Data_Type >, 130	Phylo counters, 17
count current	counter_loss
StatsCounter< Array_Type, Data_Type >, 130	Phylo counters, 17
count fun	counter_maxfuns
Counter< Array Type, Data Type >, 64	Phylo counters, 18
count_init	counter_mutual
StatsCounter< Array_Type, Data_Type >, 130	Network counters, 13
Counter	counter_neofun
Counter< Array_Type, Data_Type >, 62, 63	Phylo counters, 18
Counter< Array_Type, Data_Type >, 61	counter_neofun_a2b
~Counter, 63	Phylo counters, 18
count, 63	counter_nodecov
count_fun, 64	Network counters, 13
Counter, 62, 63	counter nodeicov
data, 65	Network counters, 13
delete data, 65	counter nodematch
desc, 65	Network counters, 13
init, 64	counter_nodeocov
init_fun, 65	Network counters, 13
name, 65	counter odegree
operator=, 64	Network counters, 14
counter_absdiff	counter_odegree15
Network counters, 11	Network counters, 14
counter_co_opt	counter_ostar2
Phylo counters, 16	Network counters, 14
counter_cogain	counter_overall_changes
Phylo counters, 16	Phylo counters, 18
counter_ctriads	counter_overall_gains
Network counters, 11	Phylo counters, 19
counter_degree	counter overall loss
Network counters, 11	Phylo counters, 19
counter density	counter_subfun
Network counters, 11	Phylo counters, 19
counter_diff	counter_ttriads
Network counters, 11	Network counters, 14
counter_edges	Counters
Network counters, 12	Counters< Array Type, Data Type >, 67
Counter_fun_type	Counters< Array Type, Data Type >, 66
typedefs.hpp, 196	~Counters, 67
COUNTER FUNCTION	add_counter, 68
barry.hpp, 156	clear, 68
counter gains	Counters, 67
Phylo counters, 16	operator=, 69
counter_gains_k_offspring	operator[], 69
Phylo counters, 17	size, 70
counter_genes_changing	Counting, 9
Phylo counters, 17	counts
counter_idegree	PhyloRuleDynData, 114
Network counters, 12	Counts_type
counter_idegree15	typedefs.hpp, 196
Network counters, 12	current_col
counter_isolates	ConstBArrayRowIter< Cell_Type, Data_Type >, 60
Network counters, 12	current_row
counter_istar2	ConstBArrayRowIter< Cell_Type, Data_Type >, 60
oountor_lotar2	Consider tray nowned $\sim$ Oen_rype, Data Type $>$ , Ot

<pre>current_stats     Support&lt; Array_Type, Data_Counter_Type,     Data_Rule_Type, Data_Rule_Dyn_Type &gt;,     140  D BArray&lt; Cell_Type, Data_Type &gt;, 32</pre>	val, 72 EXISTS, 25 AS_ONE, 25 AS_ZERO, 25 BOTH, 25 NONE, 26 ONE, 26
Rule< Array_Type, Data_Type >, 124 dat Flock, 77	TWO, 26 UKNOWN, 26
data Counter< Array_Type, Data_Type >, 65 PowerSet< Array_Type, Data_Rule_Type >, 121 default_val BArray< Cell_Type, Data_Type >, 32 delete_counters Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 141 delete_data Counter< Array_Type, Data_Type >, 65 delete_rengine Geese, 88 delete_rules Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 141 delete_rules Support< Array_Type, Data_Rule_Dyn_Type >, 141 delete_rules_dyn Support< Array_Type, Data_Counter_Type,	fill  BArrayDense < Cell_Type, Data_Type >, 53  Flock, 73  ~Flock, 74  add_data, 74  dat, 77  Flock, 74  get_counters, 74  get_support, 75  init, 75  initialized, 77  likelihood_joint, 75  nfunctions, 78  nfuns, 75  nleafs, 76  nnodes, 76  nterms, 76  ntrees, 76  operator(), 76
Data_Rule_Type, Data_Rule_Dyn_Type >, 141  delete_support	rengine, 78 set_seed, 77 support, 78
Geese, 89  desc  Counter< Array_Type, Data_Type >, 65  directed	support_size, 77  FreqTable  FreqTable < T > , 79  FrogTable < T > , 79
NetworkData, 106 duplication Node, 110 NodeData, 113 PhyloRuleDynData, 115	FreqTable < T >, 78  ~FreqTable, 79  add, 79  as_vector, 79  clear, 80  FreqTable, 79
elements_ptr BArrayDense< Cell_Type, Data_Type >, 53 elements_raw BArrayDense< Cell_Type, Data_Type >, 53 EmptyArray	get_data, 80 get_data_ptr, 80 print, 80 reserve, 80 size, 81
PowerSet< Array_Type, Data_Rule_Type >, 121 end BArrayCol< Cell_Type, Data_Type >, 47 PowerSet< Array_Type, Data_Rule_Type >, 119 Entries Entries< Cell_Type >, 71 Entries< Cell_Type >, 70 ~Entries, 71 Entries, 71 resize, 72 source, 72 target, 72	Geese, 81  ~Geese, 84  calc_reduced_sequence, 84  calc_sequence, 84  delete_rengine, 88  delete_support, 89  Geese, 83  get_counters, 84  get_probabilities, 84  get_rengine, 85  get_states, 85  get_support, 85

inherit_support, 85	get_data
init, 85	FreqTable < T >, 80
init_node, 85	PowerSet < Array_Type, Data_Rule_Type >, 119
initialized, 89	Support< Array_Type, Data_Counter_Type,
likelihood, 86	Data_Rule_Type, Data_Rule_Dyn_Type >,
likelihood_exhaust, 86	137
map_to_nodes, 89	get_data_ptr
nfunctions, 89	FreqTable < T >, 80
nfuns, 86	PowerSet < Array_Type, Data_Rule_Type >, 119
nleafs, 86	get_entries
nnodes, 86	BArray< Cell_Type, Data_Type >, 33
nodes, 89	get_last_name
nterms, 86	phylo.hpp, 172
observed_counts, 87	get_norm_const
operator=, 87	Model< Array_Type, Data_Counter_Type,
predict, 87	Data_Rule_Type, Data_Rule_Dyn_Type >,
predict_backend, 87	96
print_observed_counts, 87	get_parent
reduced_sequence, 89	Node, 109
sequence, 90	get_probabilities
set_seed, 88	Geese, 84
simulate, 88	get_pset
support_size, 88	Model< Array_Type, Data_Counter_Type,
update_annotations, 88	Data_Rule_Type, Data_Rule_Dyn_Type >,
geese-bones.hpp	96
INITIALIZED, 177	get_rengine
keygen_full, 178	Geese, 85
RULE_FUNCTION, 178	Model < Array_Type, Data_Counter_Type,
vec_diff, 178	Data_Rule_Type, Data_Rule_Dyn_Type >,
vector_caster, 178	97
get_cell BArray< Cell_Type, Data_Type >, 32	get_row BArray< Cell_Type, Data_Type >, 33
get_col	get_row_vec
BArray< Cell_Type, Data_Type >, 32	BArray< Cell_Type, Data_Type >, 33
get_col_vec	get_rules  Model< Array Type, Data Counter Type,
BArray < Cell_Type, Data_Type >, 32, 33	7= 71 / = = 71 /
get_counters	Data_Rule_Type, Data_Rule_Dyn_Type >,
Flock, 74	Support Array Time Data Counter Time
Geese, 84	Support< Array_Type, Data_Counter_Type,
Model < Array_Type, Data_Counter_Type,	Data_Rule_Type, Data_Rule_Dyn_Type >,
Data_Rule_Type, Data_Rule_Dyn_Type >,	138
96 Chata Countair & Arrest Time Data Time > 100	get_rules_dyn
StatsCounter< Array_Type, Data_Type >, 130	Model < Array_Type, Data_Counter_Type,
Support< Array_Type, Data_Counter_Type,	Data_Rule_Type, Data_Rule_Dyn_Type >,
Data_Rule_Type, Data_Rule_Dyn_Type >,	97
137	Support< Array_Type, Data_Counter_Type,
get_counts	Data_Rule_Type, Data_Rule_Dyn_Type >,
Support< Array_Type, Data_Counter_Type,	138
Data_Rule_Type, Data_Rule_Dyn_Type >,	get_seq
137	Rules < Array_Type, Data_Type >, 126
get_counts_ptr	get_states
Support< Array_Type, Data_Counter_Type,	Geese, 85
Data_Rule_Type, Data_Rule_Dyn_Type >,	get_stats
137	Model < Array_Type, Data_Counter_Type,
get_current_stats	Data_Rule_Type, Data_Rule_Dyn_Type >,
Support< Array_Type, Data_Counter_Type,	97
Data_Rule_Type, Data_Rule_Dyn_Type >,	
137	Flock, 75

Geese, 85	Counter< Array_Type, Data_Type >, 64
Model< Array_Type, Data_Counter_Type,	Flock, 75
Data_Rule_Type, Data_Rule_Dyn_Type >,	Geese, 85
97	init_fun
	Counter< Array_Type, Data_Type >, 65
id	init_node
Node, 110	Geese, 85
include/barry/barray-bones.hpp, 143	init_support
include/barry/barray-iterator.hpp, 144	PowerSet < Array_Type, Data_Rule_Type >, 119
include/barry/barray-meat-operators.hpp, 145	Support< Array_Type, Data_Counter_Type,
include/barry/barray-meat.hpp, 147	Data_Rule_Type, Data_Rule_Dyn_Type >,
include/barry/barraycell-bones.hpp, 148	138
include/barry/barraycell-meat.hpp, 149	INITIALIZED
include/barry/barraycol-bones.hpp, 150	geese-bones.hpp, 177
include/barry/barraycol-meat.hpp, 151	initialized
include/barry/barraydense-bones.hpp, 152	Flock, 77
include/barry/barry-configuration.hpp, 152	Geese, 89
include/barry/barry.hpp, 154	insert_cell
include/barry/cell-bones.hpp, 157	BArray< Cell_Type, Data_Type >, 34
include/barry/cell-meat.hpp, 157	is_empty
include/barry/col-bones.hpp, 158	BArray< Cell_Type, Data_Type >, 34
include/barry/counters-bones.hpp, 158	is_leaf
include/barry/counters-meat.hpp, 160	Node, 109
include/barry/counters/network.hpp, 161	iter
include/barry/counters/phylo.hpp, 167	ConstBArrayRowIter< Cell_Type, Data_Type >, 61
include/barry/model-bones.hpp, 173	
include/barry/model-meat.hpp, 175	keygen_default
include/barry/models/geese.hpp, 175	model-bones.hpp, 174
include/barry/models/geese/flock-bones.hpp, 176	keygen_full
include/barry/models/geese/flock-meet.hpp, 176	geese-bones.hpp, 178
include/barry/models/geese/geese-bones.hpp, 177	. lh
include/barry/models/geese/geese-meat-constructors.hpp	PhyloRuleDynData, 115
include/barry/models/geese/geese-meat-likelihood.hpp,	likelihood
179	Geese, 86
include/barry/models/geese/geese-meat-likelihood_exhau	
180	
180 include/barry/models/geese/geese-meat-predict hpp	${\sf Data\_Rule\_Type},  {\sf Data\_Rule\_Dyn\_Type} \ \ >,$
include/barry/models/geese/geese-meat-predict.hpp,	Data_Rule_Type, Data_Rule_Dyn_Type >, 98
include/barry/models/geese/geese-meat-predict.hpp, 181	Data_Rule_Type, Data_Rule_Dyn_Type >, 98 likelihood_
include/barry/models/geese/geese-meat-predict.hpp, 181 include/barry/models/geese/geese-meat-simulate.hpp,	Data_Rule_Type, Data_Rule_Dyn_Type >, 98 likelihood_ model-bones.hpp, 174
include/barry/models/geese/geese-meat-predict.hpp, 181 include/barry/models/geese/geese-meat-simulate.hpp, 181	Data_Rule_Type, Data_Rule_Dyn_Type >, 98 likelihood_ model-bones.hpp, 174 likelihood_exhaust
include/barry/models/geese/geese-meat-predict.hpp, 181 include/barry/models/geese/geese-meat-simulate.hpp, 181 include/barry/models/geese/geese-meat.hpp, 182	Data_Rule_Type, Data_Rule_Dyn_Type >, 98 likelihood_ model-bones.hpp, 174 likelihood_exhaust Geese, 86
include/barry/models/geese/geese-meat-predict.hpp, 181 include/barry/models/geese/geese-meat-simulate.hpp, 181 include/barry/models/geese/geese-meat.hpp, 182 include/barry/models/geese/geese-node-bones.hpp,	Data_Rule_Type, Data_Rule_Dyn_Type >, 98 likelihood_ model-bones.hpp, 174 likelihood_exhaust Geese, 86 likelihood_joint
include/barry/models/geese/geese-meat-predict.hpp, 181 include/barry/models/geese/geese-meat-simulate.hpp, 181 include/barry/models/geese/geese-meat.hpp, 182 include/barry/models/geese/geese-node-bones.hpp, 182	Data_Rule_Type, Data_Rule_Dyn_Type >, 98 likelihood_ model-bones.hpp, 174 likelihood_exhaust Geese, 86 likelihood_joint Flock, 75
include/barry/models/geese/geese-meat-predict.hpp, 181 include/barry/models/geese/geese-meat-simulate.hpp, 181 include/barry/models/geese/geese-meat.hpp, 182 include/barry/models/geese/geese-node-bones.hpp, 182 include/barry/powerset-bones.hpp, 183	Data_Rule_Type, Data_Rule_Dyn_Type >, 98 likelihood_ model-bones.hpp, 174 likelihood_exhaust Geese, 86 likelihood_joint Flock, 75 likelihood_total
include/barry/models/geese/geese-meat-predict.hpp,  181 include/barry/models/geese/geese-meat-simulate.hpp,  181 include/barry/models/geese/geese-meat.hpp, 182 include/barry/models/geese/geese-node-bones.hpp,  182 include/barry/powerset-bones.hpp, 183 include/barry/powerset-meat.hpp, 184	Data_Rule_Type, Data_Rule_Dyn_Type >, 98  likelihood_     model-bones.hpp, 174  likelihood_exhaust     Geese, 86  likelihood_joint     Flock, 75  likelihood_total     Model< Array_Type, Data_Counter_Type,
include/barry/models/geese/geese-meat-predict.hpp,  181 include/barry/models/geese/geese-meat-simulate.hpp,  181 include/barry/models/geese/geese-meat.hpp, 182 include/barry/models/geese/geese-node-bones.hpp,  182 include/barry/powerset-bones.hpp, 183 include/barry/powerset-meat.hpp, 184 include/barry/rules-bones.hpp, 186	Data_Rule_Type, Data_Rule_Dyn_Type >, 98 likelihood_ model-bones.hpp, 174 likelihood_exhaust Geese, 86 likelihood_joint Flock, 75 likelihood_total
include/barry/models/geese/geese-meat-predict.hpp,  181 include/barry/models/geese/geese-meat-simulate.hpp, 181 include/barry/models/geese/geese-meat.hpp, 182 include/barry/models/geese/geese-node-bones.hpp, 182 include/barry/powerset-bones.hpp, 183 include/barry/powerset-meat.hpp, 184 include/barry/rules-bones.hpp, 186 include/barry/rules-meat.hpp, 187	Data_Rule_Type, Data_Rule_Dyn_Type >, 98  likelihood_ model-bones.hpp, 174  likelihood_exhaust Geese, 86  likelihood_joint Flock, 75  likelihood_total Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,
include/barry/models/geese/geese-meat-predict.hpp, 181 include/barry/models/geese/geese-meat-simulate.hpp, 181 include/barry/models/geese/geese-meat.hpp, 182 include/barry/models/geese/geese-node-bones.hpp, 182 include/barry/powerset-bones.hpp, 183 include/barry/powerset-meat.hpp, 184 include/barry/rules-bones.hpp, 186 include/barry/rules-meat.hpp, 187 include/barry/statscounter-bones.hpp, 188	Data_Rule_Type, Data_Rule_Dyn_Type >, 98  likelihood_ model-bones.hpp, 174  likelihood_exhaust Geese, 86  likelihood_joint Flock, 75  likelihood_total Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,
include/barry/models/geese/geese-meat-predict.hpp,  181 include/barry/models/geese/geese-meat-simulate.hpp, 181 include/barry/models/geese/geese-meat.hpp, 182 include/barry/models/geese/geese-node-bones.hpp, 182 include/barry/powerset-bones.hpp, 183 include/barry/powerset-meat.hpp, 184 include/barry/rules-bones.hpp, 186 include/barry/rules-meat.hpp, 187	Data_Rule_Type, Data_Rule_Dyn_Type >, 98  likelihood_ model-bones.hpp, 174  likelihood_exhaust Geese, 86  likelihood_joint Flock, 75  likelihood_total Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 98  M PowerSet< Array_Type, Data_Rule_Type >, 121
include/barry/models/geese/geese-meat-predict.hpp, 181 include/barry/models/geese/geese-meat-simulate.hpp, 181 include/barry/models/geese/geese-meat.hpp, 182 include/barry/models/geese/geese-mode-bones.hpp, 182 include/barry/powerset-bones.hpp, 183 include/barry/powerset-meat.hpp, 184 include/barry/rules-bones.hpp, 186 include/barry/rules-meat.hpp, 187 include/barry/statscounter-bones.hpp, 188 include/barry/statscounter-meat.hpp, 189	Data_Rule_Type, Data_Rule_Dyn_Type >, 98  likelihood_ model-bones.hpp, 174  likelihood_exhaust Geese, 86  likelihood_joint Flock, 75  likelihood_total Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 98  M  PowerSet< Array_Type, Data_Rule_Type >, 121 Support< Array_Type, Data_Counter_Type,
include/barry/models/geese/geese-meat-predict.hpp,  181 include/barry/models/geese/geese-meat-simulate.hpp,  181 include/barry/models/geese/geese-meat.hpp, 182 include/barry/models/geese/geese-node-bones.hpp,  182 include/barry/powerset-bones.hpp, 183 include/barry/powerset-meat.hpp, 184 include/barry/rules-bones.hpp, 186 include/barry/rules-meat.hpp, 187 include/barry/statscounter-bones.hpp, 188 include/barry/statscounter-meat.hpp, 189 include/barry/statsdb.hpp, 190	Data_Rule_Type, Data_Rule_Dyn_Type >, 98  likelihood_ model-bones.hpp, 174  likelihood_exhaust Geese, 86  likelihood_joint Flock, 75  likelihood_total Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 98  M PowerSet< Array_Type, Data_Rule_Type >, 121
include/barry/models/geese/geese-meat-predict.hpp,  181 include/barry/models/geese/geese-meat-simulate.hpp,  181 include/barry/models/geese/geese-meat.hpp, 182 include/barry/models/geese/geese-node-bones.hpp,  182 include/barry/powerset-bones.hpp, 183 include/barry/powerset-meat.hpp, 184 include/barry/rules-bones.hpp, 186 include/barry/rules-meat.hpp, 187 include/barry/statscounter-bones.hpp, 188 include/barry/statscounter-meat.hpp, 189 include/barry/statsdb.hpp, 190 include/barry/support-bones.hpp, 191	Data_Rule_Type, Data_Rule_Dyn_Type >, 98  likelihood_ model-bones.hpp, 174  likelihood_exhaust Geese, 86  likelihood_joint Flock, 75  likelihood_total Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 98  M  PowerSet< Array_Type, Data_Rule_Type >, 121 Support< Array_Type, Data_Rule_Dyn_Type >, 141
include/barry/models/geese/geese-meat-predict.hpp,  181 include/barry/models/geese/geese-meat-simulate.hpp,  181 include/barry/models/geese/geese-meat.hpp, 182 include/barry/models/geese/geese-mode-bones.hpp,  182 include/barry/powerset-bones.hpp, 183 include/barry/powerset-meat.hpp, 184 include/barry/rules-bones.hpp, 186 include/barry/rules-meat.hpp, 187 include/barry/statscounter-bones.hpp, 188 include/barry/statscounter-meat.hpp, 189 include/barry/statsdb.hpp, 190 include/barry/support-bones.hpp, 191 include/barry/support-meat.hpp, 193	Data_Rule_Type, Data_Rule_Dyn_Type >, 98  likelihood_ model-bones.hpp, 174  likelihood_exhaust Geese, 86  likelihood_joint Flock, 75  likelihood_total Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 98  M PowerSet< Array_Type, Data_Rule_Type >, 121 Support< Array_Type, Data_Rule_Dyn_Type >, Data_Rule_Type, Data_Rule_Dyn_Type >, 141  Map
include/barry/models/geese/geese-meat-predict.hpp,  181 include/barry/models/geese/geese-meat-simulate.hpp,  181 include/barry/models/geese/geese-meat.hpp, 182 include/barry/models/geese/geese-node-bones.hpp,  182 include/barry/powerset-bones.hpp, 183 include/barry/powerset-meat.hpp, 184 include/barry/rules-bones.hpp, 186 include/barry/rules-meat.hpp, 187 include/barry/statscounter-bones.hpp, 188 include/barry/statscounter-meat.hpp, 189 include/barry/statsdb.hpp, 190 include/barry/support-bones.hpp, 191 include/barry/support-meat.hpp, 193 include/barry/typedefs.hpp, 194	Data_Rule_Type, Data_Rule_Dyn_Type >, 98  likelihood_ model-bones.hpp, 174  likelihood_exhaust Geese, 86  likelihood_joint Flock, 75  likelihood_total Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 98  M PowerSet< Array_Type, Data_Rule_Type >, 121 Support< Array_Type, Data_Rule_Dyn_Type >, 141  Map barry-configuration.hpp, 154
include/barry/models/geese/geese-meat-predict.hpp,  181 include/barry/models/geese/geese-meat-simulate.hpp, 181 include/barry/models/geese/geese-meat.hpp, 182 include/barry/models/geese/geese-node-bones.hpp, 182 include/barry/powerset-bones.hpp, 183 include/barry/powerset-meat.hpp, 184 include/barry/rules-bones.hpp, 186 include/barry/rules-meat.hpp, 187 include/barry/statscounter-bones.hpp, 188 include/barry/statscounter-meat.hpp, 189 include/barry/statsdb.hpp, 190 include/barry/support-bones.hpp, 191 include/barry/support-meat.hpp, 193 include/barry/typedefs.hpp, 194 indices	Data_Rule_Type, Data_Rule_Dyn_Type >, 98  likelihood_ model-bones.hpp, 174  likelihood_exhaust Geese, 86  likelihood_joint Flock, 75  likelihood_total Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 98  M PowerSet< Array_Type, Data_Rule_Type >, 121 Support< Array_Type, Data_Rule_Dyn_Type >, 141  Map barry-configuration.hpp, 154  map_to_nodes
include/barry/models/geese/geese-meat-predict.hpp, 181 include/barry/models/geese/geese-meat-simulate.hpp, 181 include/barry/models/geese/geese-meat.hpp, 182 include/barry/models/geese/geese-node-bones.hpp, 182 include/barry/powerset-bones.hpp, 183 include/barry/powerset-meat.hpp, 184 include/barry/rules-bones.hpp, 186 include/barry/rules-meat.hpp, 187 include/barry/statscounter-bones.hpp, 188 include/barry/statscounter-meat.hpp, 189 include/barry/statsdb.hpp, 190 include/barry/support-bones.hpp, 191 include/barry/support-meat.hpp, 193 include/barry/typedefs.hpp, 194 indices NetCounterData, 103	Data_Rule_Type, Data_Rule_Dyn_Type >, 98  likelihood_ model-bones.hpp, 174  likelihood_exhaust Geese, 86  likelihood_joint Flock, 75  likelihood_total Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 98  M PowerSet< Array_Type, Data_Rule_Type >, 121 Support< Array_Type, Data_Rule_Dyn_Type >, 141  Map barry-configuration.hpp, 154

typedefs.hpp, 196	network.hpp, 163
max_num_elements	NET_C_DATA_NUM
Support< Array_Type, Data_Counter_Type,	network.hpp, 163
Data_Rule_Type, Data_Rule_Dyn_Type >,	NetCounter
141	network.hpp, 165
Model	NetCounterData, 102
Model< Array_Type, Data_Counter_Type,	$\sim$ NetCounterData, 103
Data_Rule_Type, Data_Rule_Dyn_Type >,	indices, 103
92, 93	NetCounterData, 103
Model < Array_Type, Data_Counter_Type, Data_Rule_Type	pe, numbers, 104
Data_Rule_Dyn_Type >, 90	NetCounters
$\sim$ Model, 93	network.hpp, 165
add_array, 93	NetModel
add_counter, 94	network.hpp, 165
add_rule, 94, 95	NetRule
add_rule_dyn, 95, 96	network.hpp, 165
get_counters, 96	NetRules
get_norm_const, 96	network.hpp, 165
get_pset, 96	NetStatsCounter
get_rengine, 97	network.hpp, 165
get rules, 97	NetSupport
get_rules_dyn, 97	network.hpp, 166
get stats, 97	Network
get_support, 97	network.hpp, 166
likelihood, 98	Network counters, 10
likelihood_total, 98	counter_absdiff, 11
Model, 92, 93	counter_ctriads, 11
nterms, 99	counter_degree, 11
operator=, 99	counter_density, 11
print_stats, 99	counter_diff, 11
sample, 99, 100	counter_edges, 12
set_counters, 100	counter_idegree, 12
set_keygen, 100	counter_idegree15, 12
set_rengine, 100	counter_isolates, 12
set_rules, 101	counter_istar2, 12
set rules dyn, 101	counter mutual, 13
set_seed, 101	counter nodecov, 13
size, 101	counter_nodeicov, 13
size_unique, 101	counter_nodematch, 13
store_psets, 102	counter_nodeocov, 13
support size, 102	counter_odegree, 14
model-bones.hpp	counter_odegree15, 14
keygen_default, 174	counter_odegree 15, 14
likelihood , 174	counter_ttriads, 14
update_normalizing_constant, 174	NETWORK_COUNTER, 14
update_normalizing_constant, 174	network.hpp
N	NET_C_DATA_IDX, 163
PowerSet< Array_Type, Data_Rule_Type >, 121	
Support< Array_Type, Data_Counter_Type,	NET_C_DATA_NUM, 163
Data_Rule_Type, Data_Rule_Dyn_Type >,	NetCounter, 165
142	NetCounters, 165
name	NetRule 165
Counter< Array_Type, Data_Type >, 65	NetRule, 165
narray	NetRules, 165
Node, 110	NetStatsCounter, 165
ncol	NetSupport, 166
BArray < Cell_Type, Data_Type >, 34	Network, 166
BArrayDense < Cell_Type, Data_Type >, 53	NETWORK_COUNTER, 163
NET C DATA IDX	NETWORK_COUNTER_LAMBDA, 164

NETWORK BUILT 101	EVICTO CO
NETWORK_RULE, 164	EXISTS, 26
NETWORK_RULE_LAMBDA, 164	nrow
rules_zerodiag, 166	BArray< Cell_Type, Data_Type >, 35
NETWORK_COUNTER	BArrayDense < Cell_Type, Data_Type >, 53
Network counters, 14	nterms
network.hpp, 163	Flock, 76
NETWORK_COUNTER_LAMBDA	Geese, 86
network.hpp, 164	Model < Array_Type, Data_Counter_Type,
NETWORK RULE	Data_Rule_Type, Data_Rule_Dyn_Type >,
network.hpp, 164	99
NETWORK RULE LAMBDA	ntrees
network.hpp, 164	Flock, 76
·	
NetworkData, 104	numbers
~NetworkData, 105	NetCounterData, 104
directed, 106	observed_counts
NetworkData, 105	Geese, 87
vertex_attr, 106	
nfunctions	offspring
Flock, 78	Node, 110
Geese, 89	ONE
nfuns	CHECK, 24
Flock, 75	EXISTS, 26
Geese, 86	operator Cell_Type
nleafs	BArrayCell< Cell_Type, Data_Type >, 42
Flock, 76	BArrayCell_const< Cell_Type, Data_Type >, 45
Geese, 86	BArrayCol< Cell_Type, Data_Type >, 47
	Cell< Cell_Type >, 58
nnodes	operator!=
Flock, 76	BArrayCell_const< Cell_Type, Data_Type >, 45
Geese, 86	
nnozero	BArrayCol_const< Cell_Type, Data_Type >, 50
BArray< Cell_Type, Data_Type >, 35	operator<
Node, 106	BArrayCell_const< Cell_Type, Data_Type >, 45
$\sim$ Node, 108	BArrayCol_const< Cell_Type, Data_Type >, 50
annotations, 109	operator<=
array, 109	BArrayCell_const< Cell_Type, Data_Type >, 45
arrays, 109	BArrayCol_const< Cell_Type, Data_Type >, 50
duplication, 110	operator>
get_parent, 109	BArrayCell_const< Cell_Type, Data_Type >, 45
id, 110	BArrayCol_const< Cell_Type, Data_Type >, 51
	operator>=
is_leaf, 109	BArrayCell_const< Cell_Type, Data_Type >, 46
narray, 110	BArrayCol const< Cell Type, Data Type >, 51
Node, 107, 108	
offspring, 110	operator*=
ord, 110	BArray Cell_Type, Data_Type >, 35
parent, 111	BArrayCell < Cell_Type, Data_Type >, 42
probability, 111	BArrayCol< Cell_Type, Data_Type >, 48
subtree_prob, 111	operator()
visited, 111	BArray< Cell_Type, Data_Type >, 35
NodeData, 112	BArrayDense < Cell_Type, Data_Type >, 54
∼NodeData, 112	Flock, 76
blengths, 113	Rule < Array_Type, Data_Type >, 124
duplication, 113	Rules < Array_Type, Data_Type >, 127
•	vecHasher $<$ T $>$ , 142
NodeData, 112	operator+=
states, 113	BArray< Cell_Type, Data_Type >, 35, 36
nodes	BArrayCell< Cell_Type, Data_Type >, 42
Geese, 89	
NONE	BArrayCol < Cell_Type, Data_Type >, 48
CHECK, 24	operator-=
	BArray< Cell_Type, Data_Type >, 36

BArrayCell < Cell_Type, Data_Type >, 43	PhyloCounterData, 170
BArrayCol< Cell_Type, Data_Type >, 48	PhyloCounters, 170
operator/=	PhyloModel, 170
BArray < Cell_Type, Data_Type >, 36	PhyloPowerSet, 171
BArrayCell< Cell_Type, Data_Type >, 43	PhyloRule, 171
BArrayCol< Cell_Type, Data_Type >, 48	PhyloRuleData, 171
operator=	PhyloRuleDyn, 171
BArray< Cell_Type, Data_Type >, 37	PhyloRules, 171
BArrayCell< Cell_Type, Data_Type >, 43	PhyloRulesDyn, 171
BArrayCol< Cell_Type, Data_Type >, 48	PhyloStatsCounter, 172
Cell< Cell_Type >, 58	PhyloSupport, 172
Counter< Array_Type, Data_Type >, 64	PHYLO_CHECK_MISSING
Counters< Array_Type, Data_Type >, 69	phylo.hpp, 169
Geese, 87	PHYLO_COUNTER_LAMBDA
Model< Array_Type, Data_Counter_Type,	phylo.hpp, 169
Data_Rule_Type, Data_Rule_Dyn_Type >,	PHYLO_RULE_DYN_LAMBDA
99	phylo.hpp, 169
Rules < Array_Type, Data_Type >, 127	PhyloArray
operator==	phylo.hpp, 170
BArray< Cell_Type, Data_Type >, 37	PhyloCounter
BArrayCell< Cell_Type, Data_Type >, 43	phylo.hpp, 170
BArrayCell_const< Cell_Type, Data_Type >, 45	PhyloCounterData
BArrayCol< Cell_Type, Data_Type >, 49	phylo.hpp, 170
BArrayCol_const< Cell_Type, Data_Type >, 51	PhyloCounters
operator[]	phylo.hpp, 170
BArrayDense < Cell_Type, Data_Type >, 54	PhyloModel
Counters < Array_Type, Data_Type >, 69	phylo.hpp, 170
PowerSet< Array_Type, Data_Rule_Type >, 120	PhyloPowerSet
ord	phylo.hpp, 171
Node, 110	PhyloRule
out_of_range	phylo.hpp, 171
BArray< Cell_Type, Data_Type >, 37	PhyloRuleData
	phylo.hpp, 171
parent	PhyloRuleDyn
Node, 111	phylo.hpp, 171
Phylo counters, 15	PhyloRuleDynData, 113
counter_co_opt, 16	~PhyloRuleDynData, 114
counter_cogain, 16	counts, 114
counter_gains, 16	duplication, 115
counter_gains_k_offspring, 17	lb, 115
counter_genes_changing, 17	PhyloRuleDynData, 114
counter_longest, 17	pos, 115
counter_loss, 17	ub, 115
counter_maxfuns, 18	PhyloRules
counter_neofun, 18	phylo.hpp, 171
counter_neofun_a2b, 18	PhyloRulesDyn
counter_overall_changes, 18	phylo.hpp, 171
counter_overall_gains, 19	PhyloStatsCounter
counter_overall_loss, 19	phylo.hpp, 172
counter_subfun, 19	PhyloSupport
Phylo rules, 20	phylo.hpp, 172
rule_dyn_limit_changes, 20	pos
phylo.hpp	PhyloRuleDynData, 115
get_last_name, 172	PowerSet
PHYLO_CHECK_MISSING, 169	PowerSet < Array_Type, Data_Rule_Type >, 117
PHYLO_COUNTER_LAMBDA, 169	PowerSet < Array_Type, Data_Rule_Type >, 116
PHYLO_RULE_DYN_LAMBDA, 169	~PowerSet, 118
PhyloArray, 170	add_rule, 118
PhyloCounter, 170	444_1410, 110

begin, 118	BArray< Cell_Type, Data_Type >, 38
calc, 119	ROW
coordinates_free, 120	barray-meat-operators.hpp, 146
coordinates_locked, 120	barray-meat.hpp, 148
data, 121	row
EmptyArray, 121	BArray< Cell_Type, Data_Type >, 38
end, 119	Row_type
get_data, 119	typedefs.hpp, 196
get_data_ptr, 119	Rule
init_support, 119	Rule < Array_Type, Data_Type >, 123
M, 121	Rule < Array_Type, Data_Type >, 122
N, 121	$\sim$ Rule, 123
operator[], 120	D, 124
PowerSet, 117	operator(), 124
reset, 120	Rule, 123
rules, 121	rule_dyn_limit_changes
rules_deleted, 122	Phylo rules, 20
size, 120	rule_fun_default
predict	rules-bones.hpp, 187
Geese, 87	Rule_fun_type
predict_backend	typedefs.hpp, 197
Geese, 87	RULE_FUNCTION
print	barry.hpp, 156
BArray< Cell_Type, Data_Type >, 37	geese-bones.hpp, 178
BArrayDense < Cell_Type, Data_Type >, 54	RULE_LAMBDA
FreqTable < T >, 80	barry.hpp, 156
Support< Array_Type, Data_Counter_Type,	Rules
Data_Rule_Type, Data_Rule_Dyn_Type >,	Rules< Array_Type, Data_Type >, 125
138	rules
print_observed_counts	PowerSet< Array_Type, Data_Rule_Type >, 121
Geese, 87	Rules < Array_Type, Data_Tule_Type >, 124
print_stats	~Rules, 125
Model< Array_Type, Data_Counter_Type,	add_rule, 126
Data_Rule_Type, Data_Bule_Dyn_Type >,	
99	clear, 126
	get_seq, 126
printf_barry	operator(), 127 operator=, 127
barry-configuration.hpp, 154	•
probability	Rules, 125
Node, 111	size, 127
README.md, 198	rules-bones.hpp
reduced sequence	rule_fun_default, 187
Geese, 89	rules_deleted
rengine	PowerSet < Array_Type, Data_Rule_Type >, 122
Flock, 78	rules_zerodiag
	network.hpp, 166
reserve BArray< Cell_Type, Data_Type >, 37	
	sample
FreqTable < T >, 80	Model < Array_Type, Data_Counter_Type
reset	Data_Rule_Type, Data_Rule_Dyn_Type >
PowerSet < Array_Type, Data_Rule_Type >, 120	99, 100
reset_array	sequence
StatsCounter< Array_Type, Data_Type >, 130	Geese, 90
Support< Array_Type, Data_Counter_Type,	set_counters
Data_Rule_Type, Data_Rule_Dyn_Type >,	Model Array_Type, Data_Counter_Type
139	Data_Rule_Type, Data_Rule_Dyn_Type >
resize	100
BArray< Cell_Type, Data_Type >, 38	StatsCounter< Array_Type, Data_Type >, 131
Entries < Cell_Type >, 72	
rm_cell	

Support< Array_Type, Data_Counter_Type Data_Rule_Type, Data_Rule_Dyn_Type >	
139	get_counters, 130
set_data	reset_array, 130
BArray< Cell_Type, Data_Type >, 38	set_counters, 131
set_keygen	StatsCounter, 129
Model < Array_Type, Data_Counter_Type	e, store_psets
Data_Rule_Type, Data_Rule_Dyn_Type >	, Model< Array_Type, Data_Counter_Type,
100	Data_Rule_Type, Data_Rule_Dyn_Type >,
set_rengine	102
Model< Array_Type, Data_Counter_Type	e, subtree_prob
Data_Rule_Type, Data_Rule_Dyn_Type >	, Node, 111
100	Support
set_rules	Support< Array_Type, Data_Counter_Type,
Model< Array_Type, Data_Counter_Type	e, Data_Rule_Type, Data_Rule_Dyn_Type >,
Data_Rule_Type, Data_Rule_Dyn_Type >	, 134
101	support
Support< Array_Type, Data_Counter_Type	
Data_Rule_Type, Data_Rule_Dyn_Type >	
139	Data_Rule_Dyn_Type >, 131
set_rules_dyn	~Support, 135
Model< Array_Type, Data_Counter_Type	••
Data_Rule_Type, Data_Rule_Dyn_Type >	
101	add_rule_dyn, 136
Support< Array_Type, Data_Counter_Type	
Data_Rule_Type, Data_Rule_Dyn_Type >	
139	coordinates_free, 140
set seed	coordinates_locked, 140
Flock, 77	current_stats, 140
Geese, 88	delete_counters, 141
Model< Array_Type, Data_Counter_Type	
Data_Rule_Type, Data_Rule_Dyn_Type >	
101	get_counters, 137
simulate	get_counts, 137
Geese, 88	get_counts_ptr, 137
size	get_current_stats, 137
Counters< Array_Type, Data_Type >, 70	get_data, 137
FreqTable $\langle T \rangle$ , 81	get_rules, 138
Model< Array_Type, Data_Counter_Type	
Data Rule Type, Data Rule Dyn Type >	
101	M, 141
PowerSet< Array_Type, Data_Rule_Type >, 120	max_num_elements, 141
Rules < Array_Type, Data_Type >, 127	N, 142
size_unique	print, 138
Model< Array_Type, Data_Counter_Type	•
Data_Rule_Type, Data_Rule_Dyn_Type >	
101	set_rules, 139
source	set_rules_dyn, 139
Entries < Cell_Type >, 72	Support, 134
states	support-meat.hpp
NodeData, 113	BARRY_SUPPORT_MEAT_HPP, 194
Statistical Models, 9	support_size
StatsCounter	Flock, 77
StatsCounter StatsCounter Array_Type, Data_Type >, 129	Geese, 88
StatsCounter< Array_Type, Data_Type >, 128	Model< Array_Type, Data_Counter_Type,
~StatsCounter, 129	Data_Rule_Type, Data_Rule_Dyn_Type >,
add_counter, 129	102
count_all, 130	swap_cells
Journ an, 100	οπαρ_00113

```
{\tt BArray}{<}~{\tt Cell\_Type},~{\tt Data\_Type}>, {\color{red} \bf 39}
swap_cols
     BArray< Cell_Type, Data_Type >, 39
                                                             Node, 111
swap_rows
                                                        zero_col
    BArray< Cell_Type, Data_Type >, 39
target
                                                        zero row
     Entries < Cell_Type >, 72
toggle_cell
     BArray< Cell_Type, Data_Type >, 39
toggle_lock
     BArray< Cell_Type, Data_Type >, 39
transpose
     BArray< Cell_Type, Data_Type >, 40
TWO
     CHECK, 24
     EXISTS, 26
typedefs.hpp
     Col_type, 196
     Counter_fun_type, 196
     Counts type, 196
     MapVec type, 196
     Row_type, 196
     Rule_fun_type, 197
    uint, 197
    vec_equal, 197
    vec_equal_approx, 197
    vec_inner_prod, 198
ub
     PhyloRuleDynData, 115
uint
    typedefs.hpp, 197
UKNOWN
     EXISTS, 26
update annotations
    Geese, 88
update_normalizing_constant
     model-bones.hpp, 174
val
     Entries < Cell_Type >, 72
value
     Cell<br/>< Cell_Type >, 58
vec diff
     geese-bones.hpp, 178
vec_equal
     typedefs.hpp, 197
vec_equal_approx
     typedefs.hpp, 197
vec_inner_prod
     typedefs.hpp, 198
vecHasher< T>, 142
    operator(), 142
vector_caster
     geese-bones.hpp, 178
vertex attr
     NetworkData, 106
```

```
BArray< Cell_Type, Data_Type >, 41
Cell< Cell_Type >, 58
BArray< Cell_Type, Data_Type >, 40
BArray< Cell_Type, Data_Type >, 40
```

visited