barry: Your go-to motif accountant 0.0-1

Generated by Doxygen 1.9.1

1 Main Page	1
2 Module Index	5
2.1 Modules	5
3 Hierarchical Index	7
3.1 Class Hierarchy	
4 Class Index	9
4.1 Class List	9
5 File Index	11
5.1 File List	
6 Module Documentation	13
6.1 Counting	
6.1.1 Detailed Description	
6.1.2 Macro Definition Documentation	
6.1.2.1 IF MATCHES	
6.1.2.2 IF NOTMATCHES	
6.1.2.3 IS DUPLICATION	
6.1.2.4 IS EITHER	
<del>-</del>	
6.1.2.5 IS_SPECIATION	
6.1.2.6 MAKE_DEFM_HASHER	
6.1.2.7 MAKE_DUPL_VARS	
6.1.2.8 PHYLO_CHECK_MISSING	
6.1.2.9 PHYLO_COUNTER_LAMBDA	
6.1.2.10 PHYLO_RULE_DYN_LAMBDA	
6.1.2.11 PHYLO_RULE_LAMBDA	
6.1.3 Function Documentation	18
6.1.3.1 counter_co_opt()	18
6.1.3.2 counter_cogain()	18
6.1.3.3 counter_gains()	18
6.1.3.4 counter_gains_from_0()	19
6.1.3.5 counter_gains_k_offspring()	19
6.1.3.6 counter_genes_changing()	19
6.1.3.7 counter_k_genes_changing()	19
6.1.3.8 counter_less_than_p_prop_genes_changing()	20
6.1.3.9 counter_longest()	20
6.1.3.10 counter_loss()	20
6.1.3.11 counter_maxfuns()	20
6.1.3.12 counter_neofun()	21
6.1.3.13 counter_neofun_a2b()	21
6.1.3.14 counter_overall_changes()	21

6.1.3.15 counter_overall_gains()	21
6.1.3.16 counter_overall_gains_from_0()	22
6.1.3.17 counter_overall_loss()	22
6.1.3.18 counter_pairwise_first_gain()	22
6.1.3.19 counter_pairwise_neofun_singlefun()	22
6.1.3.20 counter_pairwise_overall_change()	23
6.1.3.21 counter_pairwise_preserving()	23
6.1.3.22 counter_preserve_pseudogene()	23
6.1.3.23 counter_prop_genes_changing()	23
6.1.3.24 counter_subfun()	24
6.1.3.25 get_last_name()	24
6.2 Statistical Models	24
6.2.1 Detailed Description	24
6.3 Network counters	24
6.3.1 Detailed Description	26
6.3.2 Function Documentation	26
6.3.2.1 counter_absdiff()	27
<b>6.3.2.2</b> counter_ctriads() [1/2]	27
6.3.2.3 counter_ctriads() [2/2]	27
6.3.2.4 counter_degree()	27
6.3.2.5 counter_density()	28
6.3.2.6 counter_diff()	28
6.3.2.7 counter_edges()	28
6.3.2.8 counter_fixed_effect()	28
<b>6.3.2.9 counter_idegree()</b> [1/2]	29
<b>6.3.2.10 counter_idegree()</b> [2/2]	29
6.3.2.11 counter_idegree15() [1/2]	29
<b>6.3.2.12 counter_idegree15()</b> [2/2]	29
<b>6.3.2.13 counter_isolates()</b> [1/2]	30
<b>6.3.2.14 counter_isolates()</b> [2/2]	30
<b>6.3.2.15</b> counter_istar2() [1/2]	30
6.3.2.16 counter_istar2() [2/2]	30
6.3.2.17 counter_logit_intercept()	30
6.3.2.18 counter_mutual()	31
6.3.2.19 counter_nodecov()	31
6.3.2.20 counter_nodeicov()	31
6.3.2.21 counter_nodematch()	31
6.3.2.22 counter_nodeocov()	31
<b>6.3.2.23</b> counter_odegree() [1/2]	32
<b>6.3.2.24 counter_odegree()</b> [2/2]	32
6.3.2.25 counter_odegree15() [1/2]	32
6.3.2.26 counter_odegree15() [2/2]	32

6.3.2.27 counter_ones()	32
<b>6.3.2.28</b> counter_ostar2() [1/2]	33
<b>6.3.2.29</b> counter_ostar2() [2/2]	33
6.3.2.30 counter_transition()	33
6.3.2.31 counter_transition_formula()	34
<b>6.3.2.32</b> counter_ttriads() [1/2]	34
<b>6.3.2.33</b> counter_ttriads() [2/2]	34
6.3.2.34 NETWORK_COUNTER()	34
6.3.2.35 rules_dont_become_zero()	35
6.3.2.36 rules_markov_fixed()	35
6.4 Phylo rules	35
6.4.1 Detailed Description	37
6.4.2 Typedef Documentation	37
6.4.2.1 DEFMCounter	37
6.4.2.2 DEFMCounters	37
6.4.2.3 DEFMModel	38
6.4.2.4 DEFMRule	38
6.4.2.5 DEFMRuleDyn	38
6.4.2.6 DEFMRules	38
6.4.2.7 DEFMRulesDyn	38
6.4.2.8 DEFMStatsCounter	38
6.4.2.9 DEFMSupport	39
6.4.3 Function Documentation	
6.4.3.1 at()	
6.4.3.2 DEFMCounterData() [1/2]	39
6.4.3.3 DEFMCounterData() [2/2]	39
6.4.3.4 DEFMData() [1/2]	39
6.4.3.5 DEFMData() [2/2]	39
<b>6.4.3.6 DEFMRuleData()</b> [1/3]	40
<b>6.4.3.7 DEFMRuleData()</b> [2/3]	40
<b>6.4.3.8 DEFMRuleData()</b> [3/3]	40
6.4.3.9 DEFMRuleDynData()	40
<b>6.4.3.10 idx()</b> [1/2]	41
<b>6.4.3.11 idx()</b> [2/2]	41
<b>6.4.3.12 is_true()</b> [1/2]	41
<b>6.4.3.13 is_true()</b> [2/2]	41
6.4.3.14 ncol()	41
6.4.3.15 nrow()	41
<b>6.4.3.16 num()</b> [1/2]	42
<b>6.4.3.17 num()</b> [2/2]	42
6.4.3.18 operator()()	42
6.4.3.19 print()	42

6.4.3.20 rule_dyn_limit_changes()	43
6.4.3.21 rule_leafs()	43
6.4.3.22 ~DEFMCounterData()	43
6.4.3.23 ∼DEFMData()	43
6.4.3.24 ∼DEFMRuleDynData()	44
6.4.4 Variable Documentation	44
6.4.4.1 array	44
6.4.4.2 counts	44
6.4.4.3 covar_sort	44
6.4.4.4 covar_used	44
6.4.4.5 covariates	45
<b>6.4.4.6 indices</b> [1/2]	45
<b>6.4.4.7 indices</b> [2/2]	45
6.4.4.8 init	45
6.4.4.9 is_motif	45
6.4.4.10 logical [1/2]	45
<b>6.4.4.11 logical</b> [2/2]	46
6.4.4.12 numbers [1/2]	46
6.4.4.13 numbers [2/2]	46
6.4.4.14 obs_start	46
6.4.4.15 X_ncol	46
6.4.4.16 X_nrow	46
6.5 Phylo counters	47
6.5.1 Detailed Description	48
6.5.2 Function Documentation	48
6.5.2.1 counter_co_opt()	48
6.5.2.2 counter_cogain()	49
6.5.2.3 counter_gains()	49
6.5.2.4 counter_gains_from_0()	49
6.5.2.5 counter_gains_k_offspring()	50
6.5.2.6 counter_genes_changing()	50
6.5.2.7 counter_k_genes_changing()	50
6.5.2.8 counter_less_than_p_prop_genes_changing()	50
6.5.2.9 counter_longest()	51
6.5.2.10 counter_loss()	51
6.5.2.11 counter_maxfuns()	51
6.5.2.12 counter_neofun()	51
6.5.2.13 counter_neofun_a2b()	52
6.5.2.14 counter_overall_changes()	52
6.5.2.15 counter_overall_gains()	52
6.5.2.16 counter_overall_gains_from_0()	52
6.5.2.17 counter_overall_loss()	53

6.5.2.18 counter_pairwise_first_gain()	. 53
6.5.2.19 counter_pairwise_neofun_singlefun()	. 53
6.5.2.20 counter_pairwise_overall_change()	. 53
6.5.2.21 counter_pairwise_preserving()	. 54
6.5.2.22 counter_preserve_pseudogene()	. 54
6.5.2.23 counter_prop_genes_changing()	. 54
6.5.2.24 counter_subfun()	. 54
7 Namespace Documentation	55
7.1 barry Namespace Reference	
7.1.1 Detailed Description	
7.2 barry::counters Namespace Reference	
7.2.1 Detailed Description	
7.3 barry::counters::network Namespace Reference	
7.4 CHECK Namespace Reference	
7.4.1 Detailed Description	
7.4.2 Variable Documentation	
7.4.2.1 BOTH	. 56
7.4.2.2 NONE	
7.4.2.3 ONE	. 56
7.4.2.4 TWO	. 57
7.5 defm Namespace Reference	. 57
7.6 EXISTS Namespace Reference	. 57
7.6.1 Detailed Description	. 57
7.6.2 Variable Documentation	. 57
7.6.2.1 AS_ONE	. 57
7.6.2.2 AS_ZERO	. 57
7.6.2.3 BOTH	. 58
7.6.2.4 NONE	. 58
7.6.2.5 ONE	. 58
7.6.2.6 TWO	. 58
7.6.2.7 UKNOWN	. 58
7.7 geese Namespace Reference	. 58
8 Class Documentation	59
8.1 BArray< Cell_Type, Data_Type > Class Template Reference	. 59
8.1.1 Detailed Description	. 61
8.1.2 Constructor & Destructor Documentation	. 62
<b>8.1.2.1 BArray()</b> [1/6]	. 62
<b>8.1.2.2 BArray()</b> [2/6]	. 62
<b>8.1.2.3 BArray()</b> [3/6]	. 62
<b>8.1.2.4 BArray()</b> [4/6]	. 63
<b>8.1.2.5 BArray()</b> [5/6]	. 63

<b>8.1.2.6 BArray()</b> [6/6]	6	33
8.1.2.7 ~BArray()	6	33
8.1.3 Member Function Documentation	ε	33
8.1.3.1 clear()	6	33
8.1.3.2 col()	6	34
8.1.3.3 D() [1/2]	6	34
<b>8.1.3.4 D()</b> [2/2]	6	34
8.1.3.5 D_ptr() [1/2]	6	34
8.1.3.6 D_ptr() [2/2]	6	34
8.1.3.7 default_val()	6	34
8.1.3.8 flush_data()	6	34
8.1.3.9 get_cell()	6	35
<b>8.1.3.10</b> get_col_vec() [1/2]	6	35
<b>8.1.3.11 get_col_vec()</b> [2/2]	6	35
8.1.3.12 get_entries()	6	35
8.1.3.13 get_row_vec() [1/2]	6	35
<b>8.1.3.14</b> get_row_vec() [2/2]	6	36
8.1.3.15 insert_cell() [1/3]	6	36
<b>8.1.3.16 insert_cell()</b> [2/3]	6	36
<b>8.1.3.17 insert_cell()</b> [3/3]	6	36
8.1.3.18 is_dense()	6	36
8.1.3.19 is_empty()	6	37
8.1.3.20 ncol()	6	37
8.1.3.21 nnozero()	6	37
8.1.3.22 nrow()	6	37
8.1.3.23 operator()() [1/2]	6	37
8.1.3.24 operator()() [2/2]	6	37
8.1.3.25 operator*=()	6	36
<b>8.1.3.26</b> operator+=() [1/3]	6	36
<b>8.1.3.27</b> operator+=() [2/3]	6	36
<b>8.1.3.28</b> operator+=() [3/3]	6	36
<b>8.1.3.29</b> operator-=() [1/3]	6	36
<b>8.1.3.30 operator-=()</b> [2/3]	6	36
<b>8.1.3.31 operator-=()</b> [3/3]	6	38
8.1.3.32 operator/=()	6	36
8.1.3.33 operator=() [1/2]	6	38
8.1.3.34 operator=() [2/2]	6	38
8.1.3.35 operator==()	6	36
8.1.3.36 out_of_range()	6	39
8.1.3.37 print()	7	70
8.1.3.38 print_n()	7	70
8.1.3.39 reserve()	7	70

8.1.3.40 resize()	70
8.1.3.41 rm_cell()	70
8.1.3.42 row()	71
8.1.3.43 set_data()	71
8.1.3.44 swap_cells()	71
8.1.3.45 swap_cols()	71
8.1.3.46 swap_rows()	72
8.1.3.47 toggle_cell()	72
8.1.3.48 toggle_lock()	72
8.1.3.49 transpose()	72
8.1.3.50 zero_col()	72
8.1.3.51 zero_row()	73
8.1.4 Friends And Related Function Documentation	73
8.1.4.1 BArrayCell< Cell_Type, Data_Type >	73
8.1.4.2 BArrayCell_const< Cell_Type, Data_Type >	73
8.1.5 Member Data Documentation	73
8.1.5.1 visited	73
8.2 BArrayCell< Cell_Type, Data_Type > Class Template Reference	74
8.2.1 Detailed Description	74
8.2.2 Constructor & Destructor Documentation	74
8.2.2.1 BArrayCell()	74
8.2.2.2 ~BArrayCell()	74
8.2.3 Member Function Documentation	75
8.2.3.1 operator Cell_Type()	75
8.2.3.2 operator*=()	75
8.2.3.3 operator+=()	75
8.2.3.4 operator-=()	75
8.2.3.5 operator/=()	75
8.2.3.6 operator=()	76
8.2.3.7 operator==()	76
8.3 BArrayCell_const< Cell_Type, Data_Type > Class Template Reference	76
8.3.1 Detailed Description	76
8.3.2 Constructor & Destructor Documentation	77
8.3.2.1 BArrayCell_const()	77
8.3.2.2 ~BArrayCell_const()	77
8.3.3 Member Function Documentation	77
8.3.3.1 operator Cell_Type()	77
8.3.3.2 operator"!=()	77
8.3.3.3 operator<()	78
8.3.3.4 operator<=()	78
8.3.3.5 operator==()	78
8.3.3.6 operator>()	78

8.3.3.7 operator>=()	 78
8.4 BArrayDense < Cell_Type, Data_Type > Class Template Reference	 79
8.4.1 Detailed Description	 81
8.4.2 Constructor & Destructor Documentation	 82
8.4.2.1 BArrayDense() [1/6]	 82
<b>8.4.2.2 BArrayDense()</b> [2/6]	 82
<b>8.4.2.3 BArrayDense()</b> [3/6]	 82
<b>8.4.2.4 BArrayDense()</b> [4/6]	 83
<b>8.4.2.5 BArrayDense()</b> [5/6]	 83
<b>8.4.2.6 BArrayDense()</b> [6/6]	 83
8.4.2.7 ~BArrayDense()	 83
8.4.3 Member Function Documentation	 83
8.4.3.1 clear()	 84
8.4.3.2 col() [1/2]	 84
<b>8.4.3.3 col()</b> [2/2]	 84
8.4.3.4 colsum()	 84
<b>8.4.3.5 D()</b> [1/2]	 84
<b>8.4.3.6 D()</b> [2/2]	 85
8.4.3.7 D_ptr() [1/2]	 85
8.4.3.8 D_ptr() [2/2]	 85
8.4.3.9 default_val()	 85
8.4.3.10 get_cell()	 85
8.4.3.11 get_col_vec() [1/2]	 86
<b>8.4.3.12 get_col_vec()</b> [2/2]	 86
8.4.3.13 get_data()	 86
8.4.3.14 get_entries()	 86
8.4.3.15 get_row_vec() [1/2]	 87
8.4.3.16 get_row_vec() [2/2]	 87
8.4.3.17 insert_cell() [1/2]	 87
8.4.3.18 insert_cell() [2/2]	 87
8.4.3.19 is_dense()	 88
8.4.3.20 is_empty()	 88
8.4.3.21 ncol()	 88
8.4.3.22 nnozero()	 88
8.4.3.23 nrow()	 88
8.4.3.24 operator()() [1/2]	 89
<b>8.4.3.25</b> operator()() [2/2]	 89
8.4.3.26 operator*=()	 89
<b>8.4.3.27 operator+=()</b> [1/3]	 89
<b>8.4.3.28 operator+=()</b> [2/3]	 89
<b>8.4.3.29 operator+=()</b> [3/3]	 90
8.4.3.30 operator-=() [1/3]	 90

8.4.3.31 operator-=() [2/3]	90
8.4.3.32 operator-=() [3/3]	90
8.4.3.33 operator/=()	90
8.4.3.34 operator=() [1/2]	91
<b>8.4.3.35 operator=()</b> [2/2]	91
8.4.3.36 operator==()	91
8.4.3.37 out_of_range()	91
8.4.3.38 print()	91
8.4.3.39 reserve()	92
8.4.3.40 resize()	92
8.4.3.41 rm_cell()	92
<b>8.4.3.42 row()</b> [1/2]	92
<b>8.4.3.43 row()</b> [2/2]	92
8.4.3.44 rowsum()	93
8.4.3.45 set_data()	93
8.4.3.46 swap_cells()	93
8.4.3.47 swap_cols()	93
8.4.3.48 swap_rows()	94
8.4.3.49 toggle_cell()	94
8.4.3.50 toggle_lock()	94
8.4.3.51 transpose()	94
8.4.3.52 zero_col()	94
8.4.3.53 zero_row()	95
8.4.4 Friends And Related Function Documentation	95
8.4.4.1 BArrayDenseCell< Cell_Type, Data_Type >	95
8.4.4.2 BArrayDenseCol< Cell_Type, Data_Type >	95
8.4.4.3 BArrayDenseCol_const< Cell_Type, Data_Type >	95
8.4.4.4 BArrayDenseRow< Cell_Type, Data_Type >	95
8.4.4.5 BArrayDenseRow_const< Cell_Type, Data_Type >	96
8.4.5 Member Data Documentation	96
8.4.5.1 visited	96
8.5 BArrayDenseCell< Cell_Type, Data_Type > Class Template Reference	96
8.5.1 Detailed Description	97
8.5.2 Constructor & Destructor Documentation	97
8.5.2.1 BArrayDenseCell()	97
8.5.2.2 ~BArrayDenseCell()	97
8.5.3 Member Function Documentation	97
8.5.3.1 operator Cell_Type()	97
8.5.3.2 operator*=()	98
8.5.3.3 operator+=()	98
8.5.3.4 operator-=()	98
8.5.3.5 operator/=()	98

8.5.3.6 operator=() [1/2]	98
8.5.3.7 operator=() [2/2]	99
8.5.3.8 operator==()	99
8.5.4 Friends And Related Function Documentation	99
8.5.4.1 BArrayDense < Cell_Type, Data_Type >	99
8.5.4.2 BArrayDenseCol< Cell_Type, Data_Type >	99
8.5.4.3 BArrayDenseCol_const< Cell_Type, Data_Type >	99
8.6 BArrayDenseCell_const< Cell_Type, Data_Type > Class Template Reference	100
8.6.1 Detailed Description	100
8.7 BArrayDenseCol< Cell_Type, Data_Type > Class Template Reference	100
8.7.1 Detailed Description	100
8.7.2 Constructor & Destructor Documentation	100
8.7.2.1 BArrayDenseCol()	101
8.7.3 Member Function Documentation	101
8.7.3.1 begin()	101
8.7.3.2 end()	101
8.7.3.3 operator()()	101
8.7.3.4 size()	101
8.7.4 Friends And Related Function Documentation	102
8.7.4.1 BArrayDense < Cell_Type, Data_Type >	102
8.7.4.2 BArrayDenseCell< Cell_Type, Data_Type >	102
8.7.4.3 BArrayDenseCell_const< Cell_Type, Data_Type >	102
$8.8 \; BArrayDenseCol\_const < Cell\_Type, \; Data\_Type > Class \; Template \; Reference  \dots  \dots  \dots$	102
8.8.1 Detailed Description	103
8.8.2 Constructor & Destructor Documentation	103
8.8.2.1 BArrayDenseCol_const()	103
8.8.3 Member Function Documentation	103
8.8.3.1 begin()	103
8.8.3.2 end()	103
8.8.3.3 operator()()	104
8.8.3.4 size()	104
8.8.4 Friends And Related Function Documentation	104
8.8.4.1 BArrayDenseCell< Cell_Type, Data_Type >	104
8.8.4.2 BArrayDenseCell_const< Cell_Type, Data_Type >	104
8.9 BArrayDenseRow< Cell_Type, Data_Type > Class Template Reference	104
8.9.1 Detailed Description	105
8.9.2 Constructor & Destructor Documentation	105
8.9.2.1 BArrayDenseRow()	105
8.9.3 Member Function Documentation	105
8.9.3.1 begin()	105
8.9.3.2 end()	106
8.9.3.3 operator()()	106

8.9.3.4 size()	106
8.9.4 Friends And Related Function Documentation	106
8.9.4.1 BArrayDense < Cell_Type, Data_Type >	106
8.9.4.2 BArrayDenseCell< Cell_Type, Data_Type >	106
8.9.4.3 BArrayDenseCell_const< Cell_Type, Data_Type >	107
8.10 BArrayDenseRow_const< Cell_Type, Data_Type > Class Template Reference	107
8.10.1 Detailed Description	107
8.10.2 Constructor & Destructor Documentation	107
8.10.2.1 BArrayDenseRow_const()	108
8.10.3 Member Function Documentation	108
8.10.3.1 begin()	108
8.10.3.2 end()	108
8.10.3.3 operator()()	108
8.10.3.4 size()	108
8.10.4 Friends And Related Function Documentation	109
8.10.4.1 BArrayDenseCell< Cell_Type, Data_Type >	109
8.10.4.2 BArrayDenseCell_const< Cell_Type, Data_Type >	109
8.11 BArrayRow< Cell_Type, Data_Type > Class Template Reference	109
8.11.1 Detailed Description	109
8.11.2 Constructor & Destructor Documentation	110
8.11.2.1 BArrayRow()	110
8.11.2.2 ~BArrayRow()	110
8.11.3 Member Function Documentation	110
8.11.3.1 operator BArrayRow< Cell_Type, Data_Type >()	110
8.11.3.2 operator*=()	110
8.11.3.3 operator+=()	110
8.11.3.4 operator-=()	111
8.11.3.5 operator/=()	111
8.11.3.6 operator=()	111
8.11.3.7 operator==()	111
8.12 BArrayRow_const< Cell_Type, Data_Type > Class Template Reference	111
8.12.1 Detailed Description	112
8.12.2 Constructor & Destructor Documentation	112
8.12.2.1 BArrayRow_const()	112
8.12.2.2 ~BArrayRow_const()	112
8.12.3 Member Function Documentation	112
8.12.3.1 operator BArrayRow_const< Cell_Type, Data_Type >()	112
8.12.3.2 operator"!=()	112
8.12.3.3 operator<()	113
8.12.3.4 operator<=()	113
8.12.3.5 operator==()	113
8.12.3.6 operator>()	113

8.12.3.7 operator>=()	. 113
8.13 BArrayVector< Cell_Type, Data_Type > Class Template Reference	. 113
8.13.1 Detailed Description	. 114
8.13.2 Constructor & Destructor Documentation	. 114
8.13.2.1 BArrayVector()	. 114
8.13.2.2 ~BArrayVector()	. 115
8.13.3 Member Function Documentation	. 115
8.13.3.1 begin()	. 115
8.13.3.2 end()	. 115
8.13.3.3 is_col()	. 115
8.13.3.4 is_row()	. 116
8.13.3.5 operator std::vector< Cell_Type >()	. 116
8.13.3.6 operator*=()	. 116
8.13.3.7 operator+=()	. 116
8.13.3.8 operator-=()	. 116
8.13.3.9 operator/=()	. 117
8.13.3.10 operator=()	. 117
8.13.3.11 operator==()	. 117
8.13.3.12 size()	. 117
8.14 BArrayVector_const< Cell_Type, Data_Type > Class Template Reference	. 117
8.14.1 Detailed Description	. 118
8.14.2 Constructor & Destructor Documentation	. 118
8.14.2.1 BArrayVector_const()	. 118
8.14.2.2 ~BArrayVector_const()	. 118
8.14.3 Member Function Documentation	. 119
8.14.3.1 begin()	. 119
8.14.3.2 end()	. 119
8.14.3.3 is_col()	. 119
8.14.3.4 is_row()	. 119
8.14.3.5 operator std::vector< Cell_Type >()	. 119
8.14.3.6 operator"!=()	. 120
8.14.3.7 operator<()	. 120
8.14.3.8 operator<=()	. 120
8.14.3.9 operator==()	. 120
8.14.3.10 operator>()	. 120
8.14.3.11 operator>=()	. 121
8.14.3.12 size()	. 121
8.15 Cell < Cell_Type > Class Template Reference	. 121
8.15.1 Detailed Description	. 122
8.15.2 Constructor & Destructor Documentation	. 122
8.15.2.1 Cell() [1/7]	. 122
8.15.2.2 Cell() [2/7]	. 122

8.15.2.3 ~Cell()	122
8.15.2.4 Cell() [3/7]	123
8.15.2.5 Cell() [4/7]	123
<b>8.15.2.6 Cell()</b> [5/7]	123
8.15.2.7 Cell() [6/7]	123
8.15.2.8 Cell() [7/7]	123
8.15.3 Member Function Documentation	123
8.15.3.1 add() [1/4]	124
<b>8.15.3.2 add()</b> [2/4]	124
<b>8.15.3.3 add()</b> [3/4]	124
8.15.3.4 add() [4/4]	124
8.15.3.5 operator Cell_Type()	124
8.15.3.6 operator"!=()	124
8.15.3.7 operator=() [1/2]	125
8.15.3.8 operator=() [2/2]	125
8.15.3.9 operator==()	125
8.15.4 Member Data Documentation	125
8.15.4.1 active	125
8.15.4.2 value	125
8.15.4.3 visited	126
8.16 Cell_const< Cell_Type > Class Template Reference	126
8.16.1 Detailed Description	126
8.17 ConstBArrayRowIter< Cell_Type, Data_Type > Class Template Reference	126
8.17.1 Detailed Description	127
8.17.2 Constructor & Destructor Documentation	127
8.17.2.1 ConstBArrayRowlter()	127
8.17.2.2 ~ConstBArrayRowIter()	127
8.17.3 Member Data Documentation	127
8.17.3.1 Array	128
8.17.3.2 current_col	128
8.17.3.3 current_row	128
8.17.3.4 iter	128
8.18 Counter< Array_Type, Data_Type > Class Template Reference	128
8.18.1 Detailed Description	129
8.18.2 Constructor & Destructor Documentation	130
8.18.2.1 Counter() [1/4]	130
8.18.2.2 Counter() [2/4]	130
8.18.2.3 Counter() [3/4]	130
8.18.2.4 Counter() [4/4]	130
8.18.2.5 ~Counter()	131
8.18.3 Member Function Documentation	131
8.18.3.1 count()	131

8.18.3.2 get_description()	 131
8.18.3.3 get_hasher()	 131
8.18.3.4 get_name()	 131
8.18.3.5 init()	 131
8.18.3.6 operator=() [1/2]	 132
8.18.3.7 operator=() [2/2]	 132
8.18.3.8 set_hasher()	 132
8.18.4 Member Data Documentation	 132
8.18.4.1 count_fun	 132
8.18.4.2 data	 133
8.18.4.3 desc	 133
8.18.4.4 hasher_fun	 133
8.18.4.5 init_fun	 133
8.18.4.6 name	 133
8.19 Counters< Array_Type, Data_Type > Class Template Reference	 134
8.19.1 Detailed Description	 134
8.19.2 Constructor & Destructor Documentation	 134
8.19.2.1 Counters() [1/3]	 135
8.19.2.2 ~Counters()	 135
<b>8.19.2.3 Counters()</b> [2/3]	 135
<b>8.19.2.4 Counters()</b> [3/3]	 135
8.19.3 Member Function Documentation	 135
<b>8.19.3.1 add_counter()</b> [1/2]	 136
<b>8.19.3.2</b> add_counter() [2/2]	 136
8.19.3.3 add_hash()	 136
8.19.3.4 gen_hash()	 136
8.19.3.5 get_descriptions()	 137
8.19.3.6 get_names()	 137
8.19.3.7 operator=() [1/2]	 137
<b>8.19.3.8 operator=()</b> [2/2]	 137
8.19.3.9 operator[]()	 138
8.19.3.10 size()	 138
8.20 DEFM Class Reference	 139
8.20.1 Detailed Description	 140
8.20.2 Constructor & Destructor Documentation	 140
8.20.2.1 DEFM()	 140
8.20.3 Member Function Documentation	 140
8.20.3.1 get_column_major()	 140
8.20.3.2 get_ID()	 140
8.20.3.3 get_m_order()	 141
8.20.3.4 get_model()	 141
8.20.3.5 get_n_covars()	 141

8.20.3.6 get_n_obs()	141
8.20.3.7 get_n_rows()	141
8.20.3.8 get_n_y()	141
8.20.3.9 get_X()	142
8.20.3.10 get_X_names()	142
8.20.3.11 get_Y()	142
8.20.3.12 get_Y_names()	142
8.20.3.13 init()	142
8.20.3.14 is_motif()	142
8.20.3.15 logodds()	143
8.20.3.16 motif_census()	143
8.20.3.17 print()	143
8.20.3.18 set_names()	143
8.20.3.19 simulate()	143
8.21 DEFMCounterData Class Reference	144
8.21.1 Detailed Description	144
8.22 DEFMData Class Reference	144
8.22.1 Detailed Description	145
8.23 DEFMRuleData Class Reference	146
8.23.1 Detailed Description	146
8.24 DEFMRuleDynData Class Reference	147
8.24.1 Detailed Description	147
8.25 Entries < Cell_Type > Class Template Reference	148
8.25.1 Detailed Description	148
8.25.2 Constructor & Destructor Documentation	148
8.25.2.1 Entries() [1/2]	148
8.25.2.2 Entries() [2/2]	149
8.25.2.3 ~Entries()	149
8.25.3 Member Function Documentation	149
8.25.3.1 resize()	149
8.25.4 Member Data Documentation	149
8.25.4.1 source	149
8.25.4.2 target	149
8.25.4.3 val	150
8.26 Flock Class Reference	150
8.26.1 Detailed Description	151
8.26.2 Constructor & Destructor Documentation	151
8.26.2.1 Flock()	151
8.26.2.2 ~Flock()	151
8.26.3 Member Function Documentation	151
8.26.3.1 add_data()	151
8.26.3.2 colnames()	152

8.26.3.3 get_counters()	 152
8.26.3.4 get_model()	 152
8.26.3.5 get_stats_support()	 152
8.26.3.6 get_stats_target()	 153
8.26.3.7 get_support_fun()	 153
8.26.3.8 init()	 153
8.26.3.9 likelihood_joint()	 153
8.26.3.10 nfuns()	 154
8.26.3.11 nleafs()	 154
8.26.3.12 nnodes()	 154
8.26.3.13 nterms()	 154
8.26.3.14 ntrees()	 154
8.26.3.15 operator()()	 154
8.26.3.16 parse_polytomies()	 155
8.26.3.17 print()	 155
8.26.3.18 set_seed()	 155
8.26.3.19 support_size()	 156
8.26.4 Member Data Documentation	 156
8.26.4.1 dat	 156
8.26.4.2 initialized	 156
8.26.4.3 model	 156
8.26.4.4 nfunctions	 156
8.26.4.5 rengine	 157
8.27 FreqTable < T > Class Template Reference	 157
8.27.1 Detailed Description	 157
8.27.2 Constructor & Destructor Documentation	 158
8.27.2.1 FreqTable()	 158
8.27.2.2 ∼FreqTable()	 158
8.27.3 Member Function Documentation	 158
8.27.3.1 add()	 158
8.27.3.2 as_vector()	 158
8.27.3.3 clear()	 159
8.27.3.4 get_data()	 159
8.27.3.5 get_index()	 159
8.27.3.6 make_hash()	 159
8.27.3.7 print()	 159
8.27.3.8 reserve()	 160
8.27.3.9 size()	 160
8.28 Geese Class Reference	 160
8.28.1 Detailed Description	 163
8.28.2 Constructor & Destructor Documentation	 163
8.28.2.1 Geese() [1/4]	 164

8.28.2.2 Geese() [2/4]	 164
<b>8.28.2.3 Geese()</b> [3/4]	 164
8.28.2.4 Geese() [4/4]	 164
8.28.2.5 ~Geese()	 164
8.28.3 Member Function Documentation	 164
8.28.3.1 calc_reduced_sequence()	 165
8.28.3.2 calc_sequence()	 165
8.28.3.3 colnames()	 165
8.28.3.4 get_annotated_nodes()	 165
8.28.3.5 get_annotations()	 165
8.28.3.6 get_counters()	 166
8.28.3.7 get_model()	 166
8.28.3.8 get_probabilities()	 166
8.28.3.9 get_rengine()	 166
8.28.3.10 get_states()	 166
8.28.3.11 get_support_fun()	 167
8.28.3.12 inherit_support()	 167
8.28.3.13 init()	 167
8.28.3.14 init_node()	 167
8.28.3.15 likelihood()	 167
8.28.3.16 likelihood_exhaust()	 168
8.28.3.17 nannotations()	 168
8.28.3.18 nfuns()	 168
8.28.3.19 nleafs()	 168
8.28.3.20 nnodes()	 168
8.28.3.21 nterms()	 169
8.28.3.22 observed_counts()	 169
8.28.3.23 operator=() [1/2]	 169
8.28.3.24 operator=() [2/2]	 169
8.28.3.25 parse_polytomies()	 169
8.28.3.26 predict()	 170
8.28.3.27 predict_backend()	 170
8.28.3.28 predict_exhaust()	 170
8.28.3.29 predict_exhaust_backend()	 170
8.28.3.30 predict_sim()	 170
8.28.3.31 print()	 171
8.28.3.32 print_nodes()	 171
8.28.3.33 print_observed_counts()	 171
8.28.3.34 set_seed()	 171
8.28.3.35 simulate()	 171
8.28.3.36 support_size()	 171
8.28.3.37 update_annotations()	 172

8.28.4 Member Data Documentation		172
8.28.4.1 delete_rengine		172
8.28.4.2 delete_support		172
8.28.4.3 etype_default		172
8.28.4.4 etype_duplication		172
8.28.4.5 etype_either		173
8.28.4.6 etype_speciation		173
8.28.4.7 initialized		173
8.28.4.8 map_to_state_id		173
8.28.4.9 nfunctions		173
8.28.4.10 nodes		173
8.28.4.11 pset_loc		174
8.28.4.12 reduced_sequence		174
8.28.4.13 sequence		174
8.29 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class Te	m-	
plate Reference		
8.29.1 Detailed Description		179
8.29.2 Constructor & Destructor Documentation		179
<b>8.29.2.1 Model()</b> [1/3]		180
<b>8.29.2.2 Model()</b> [2/3]		180
<b>8.29.2.3 Model()</b> [3/3]		180
8.29.2.4 ~ Model()		180
8.29.3 Member Function Documentation		180
8.29.3.1 add_array()		180
<b>8.29.3.2</b> add_counter() [1/2]		181
<b>8.29.3.3 add_counter()</b> [2/2]		181
8.29.3.4 add_hasher()		181
<b>8.29.3.5</b> add_rule() [1/2]		182
<b>8.29.3.6 add_rule()</b> [2/2]		182
<b>8.29.3.7 add_rule_dyn()</b> [1/2]		182
<b>8.29.3.8 add_rule_dyn()</b> [2/2]		182
8.29.3.9 colnames()		182
8.29.3.10 conditional_prob()		183
8.29.3.11 gen_key()		183
8.29.3.12 get_arrays2support()		183
8.29.3.13 get_counters()		184
8.29.3.14 get_likelihoods()		184
8.29.3.15 get_normalizing_constants()		184
8.29.3.16 get_pset()		184
8.29.3.17 get_pset_arrays()		184
8.29.3.18 get_pset_locations()		185
8.29.3.19 get_pset_probs()		185

	8.29.3.20 get_pset_sizes()	15
	8.29.3.21 get_pset_stats() [1/2]	35
	8.29.3.22 get_pset_stats() [2/2]	36
	8.29.3.23 get_rengine()	36
	8.29.3.24 get_rules()	6
	8.29.3.25 get_rules_dyn()	6
	8.29.3.26 get_stats_support()	6
	8.29.3.27 get_stats_support_sizes()	37
	8.29.3.28 get_stats_support_sizes_acc()	37
	8.29.3.29 get_stats_target()	37
	8.29.3.30 get_support_fun()	37
	8.29.3.31 likelihood() [1/4]	8
	8.29.3.32 likelihood() [2/4]	8
	8.29.3.33 likelihood() [3/4]	8
	8.29.3.34 likelihood() [4/4]	19
	8.29.3.35 likelihood_total()	19
	8.29.3.36 nrules()	19
	8.29.3.37 nrules_dyn()	9
	8.29.3.38 nterms()	0
	8.29.3.39 operator=()	0
	8.29.3.40 print()	0
	8.29.3.41 print_stats()	0
	8.29.3.42 sample() [1/2]	1
	8.29.3.43 sample() [2/2]	1
	8.29.3.44 set_counters()	)1
	8.29.3.45 set_rengine()	)1
	8.29.3.46 set_rules()	)2
	8.29.3.47 set_rules_dyn()	)2
	8.29.3.48 set_seed()	)2
	8.29.3.49 set_transform_model()	)2
	8.29.3.50 size()	)3
	8.29.3.51 size_unique()	)3
	8.29.3.52 store_psets()	)3
	8.29.3.53 support_size()	)3
	8.29.3.54 transform_model()	)4
	8.29.3.55 update_likelihoods()	)4
	8.29.3.56 update_normalizing_constants()	)4
	8.29.3.57 update_pset_probs()	)4
8.29.4 N	lember Data Documentation	)5
	8.29.4.1 arrays2support	
	8.29.4.2 counter_fun	)5
	8.29.4.3 counters	15

8.29.4.4 delete_counters	195
8.29.4.5 delete_rengine	196
8.29.4.6 delete_rules	196
8.29.4.7 delete_rules_dyn	196
8.29.4.8 first_calc_done	196
8.29.4.9 keys2support	196
8.29.4.10 normalizing_constants	197
8.29.4.11 params_last	197
8.29.4.12 pset_arrays	197
8.29.4.13 pset_locations	197
8.29.4.14 pset_probs	198
8.29.4.15 pset_sizes	198
8.29.4.16 pset_stats	198
8.29.4.17 rengine	198
8.29.4.18 rules	199
8.29.4.19 rules_dyn	199
8.29.4.20 stats_likelihood	199
8.29.4.21 stats_support	199
8.29.4.22 stats_support_n_arrays	200
8.29.4.23 stats_support_sizes	200
8.29.4.24 stats_support_sizes_acc	200
8.29.4.25 stats_target	200
8.29.4.26 support_fun	201
8.29.4.27 transform_model_fun	201
8.29.4.28 transform_model_term_names	201
8.29.4.29 with_pset	202
8.30 NetCounterData Class Reference	202
8.30.1 Detailed Description	202
8.30.2 Constructor & Destructor Documentation	202
<b>8.30.2.1 NetCounterData()</b> [1/2]	202
<b>8.30.2.2 NetCounterData()</b> [2/2]	203
8.30.2.3 ~NetCounterData()	203
8.30.3 Member Data Documentation	203
8.30.3.1 indices	203
8.30.3.2 numbers	203
8.31 NetworkData Class Reference	203
8.31.1 Detailed Description	204
8.31.2 Constructor & Destructor Documentation	204
<b>8.31.2.1 NetworkData()</b> [1/3]	204
<b>8.31.2.2</b> NetworkData() [2/3]	204
<b>8.31.2.3 NetworkData()</b> [3/3]	205
8.31.2.4 ~NetworkData()	205

8.31.3 Member Data Documentation	)5
8.31.3.1 directed	)5
8.31.3.2 vertex_attr	ე6
8.32 Node Class Reference	ე6
8.32.1 Detailed Description	)7
8.32.2 Constructor & Destructor Documentation	)7
8.32.2.1 Node() [1/5]	)7
8.32.2.2 Node() [2/5]	38
8.32.2.3 Node() [3/5]	38
8.32.2.4 Node() [4/5]	38
8.32.2.5 Node() [5/5]	38
8.32.2.6 ∼Node()	38
8.32.3 Member Function Documentation	38
8.32.3.1 get_parent()	)9
8.32.3.2 is_leaf()	)9
8.32.3.3 noffspring()	)9
8.32.4 Member Data Documentation	)9
8.32.4.1 annotations	)9
8.32.4.2 array	)9
8.32.4.3 arrays	
8.32.4.4 duplication	10
8.32.4.5 id	10
8.32.4.6 narray	10
8.32.4.7 offspring	10
8.32.4.8 ord	11
8.32.4.9 parent	11
8.32.4.10 probability	11
8.32.4.11 subtree_prob	11
8.32.4.12 visited	11
8.33 NodeData Class Reference	
8.33.1 Detailed Description	
8.33.2 Constructor & Destructor Documentation	12
8.33.2.1 NodeData()	
8.33.3 Member Data Documentation	
8.33.3.1 blengths	13
8.33.3.2 duplication	13
8.33.3.3 has_leaf	13
8.33.3.4 states	
8.34 PhyloCounterData Class Reference	
8.34.1 Detailed Description	
8.34.2 Constructor & Destructor Documentation	
8.34.2.1 PhyloCounterData() [1/2]	14

<b>8.34.2.2</b> PhyloCounterData() [2/2]	214
8.34.3 Member Function Documentation	214
8.34.3.1 at()	215
8.34.3.2 begin()	215
8.34.3.3 empty()	215
8.34.3.4 end()	215
8.34.3.5 get_counters()	215
8.34.3.6 operator()()	215
8.34.3.7 operator[]()	216
8.34.3.8 push_back()	216
8.34.3.9 reserve()	216
8.34.3.10 shrink_to_fit()	216
8.34.3.11 size()	216
8.35 PhyloRuleDynData Class Reference	216
8.35.1 Detailed Description	217
8.35.2 Constructor & Destructor Documentation	217
8.35.2.1 PhyloRuleDynData()	217
8.35.2.2 $\sim$ PhyloRuleDynData()	217
8.35.3 Member Function Documentation	217
8.35.3.1 operator()()	218
8.35.4 Member Data Documentation	218
8.35.4.1 counts	218
8.35.4.2 duplication	218
8.35.4.3 lb	218
8.35.4.4 pos	218
8.35.4.5 ub	218
8.36 PowerSet< Array_Type, Data_Rule_Type > Class Template Reference	219
8.36.1 Detailed Description	220
8.36.2 Constructor & Destructor Documentation	220
<b>8.36.2.1 PowerSet()</b> [1/3]	220
<b>8.36.2.2 PowerSet()</b> [2/3]	220
<b>8.36.2.3 PowerSet()</b> [3/3]	221
8.36.2.4 ~PowerSet()	221
8.36.3 Member Function Documentation	221
8.36.3.1 add_rule() [1/2]	221
<b>8.36.3.2 add_rule()</b> [2/2]	221
8.36.3.3 begin()	221
8.36.3.4 calc()	222
8.36.3.5 end()	222
8.36.3.6 get_data()	222
8.36.3.7 get_data_ptr()	222
8.36.3.8 init_support()	222

8.36.3.9 operator[]()	23
8.36.3.10 reset()	23
8.36.3.11 size()	23
8.36.4 Member Data Documentation	23
8.36.4.1 coordinates_free	23
8.36.4.2 coordinates_locked	23
8.36.4.3 data	24
8.36.4.4 EmptyArray	24
8.36.4.5 M	24
8.36.4.6 N	24
8.36.4.7 n_free	24
8.36.4.8 n_locked	24
8.36.4.9 rules	25
8.36.4.10 rules_deleted	25
8.37 Progress Class Reference	25
8.37.1 Detailed Description	25
8.37.2 Constructor & Destructor Documentation	25
8.37.2.1 Progress()	26
8.37.2.2 ~Progress()	26
8.37.3 Member Function Documentation	26
8.37.3.1 end()	26
8.37.3.2 next()	26
8.38 Rule < Array_Type, Data_Type > Class Template Reference	26
8.38.1 Detailed Description	27
8.38.2 Constructor & Destructor Documentation	27
8.38.2.1 Rule() [1/2]	27
8.38.2.2 Rule() [2/2]	27
8.38.2.3 ~Rule()	28
8.38.3 Member Function Documentation	28
8.38.3.1 D()	28
8.38.3.2 get_description() [1/2]	28
8.38.3.3 get_description() [2/2]	28
8.38.3.4 get_name() [1/2]	28
8.38.3.5 get_name() [2/2]	29
8.38.3.6 operator()()	29
8.39 Rules < Array_Type, Data_Type > Class Template Reference	29
8.39.1 Detailed Description	30
8.39.2 Constructor & Destructor Documentation	30
8.39.2.1 Rules() [1/2]	30
8.39.2.2 Rules() [2/2]	30
8.39.2.3 ~Rules()	31
8 39 3 Member Function Documentation	21

8.39.3.1 add_rule() [1/2]	231
<b>8.39.3.2</b> add_rule() [2/2]	231
8.39.3.3 begin()	231
8.39.3.4 end()	231
8.39.3.5 get_descriptions()	232
8.39.3.6 get_names()	232
8.39.3.7 get_seq()	232
8.39.3.8 operator()()	232
8.39.3.9 operator=()	233
8.39.3.10 size()	233
8.40 StatsCounter< Array_Type, Data_Type > Class Template Reference	233
8.40.1 Detailed Description	234
8.40.2 Constructor & Destructor Documentation	234
8.40.2.1 StatsCounter() [1/3]	234
<b>8.40.2.2 StatsCounter()</b> [2/3]	235
<b>8.40.2.3 StatsCounter()</b> [3/3]	235
8.40.2.4 ~StatsCounter()	235
8.40.3 Member Function Documentation	235
8.40.3.1 add_counter()	235
8.40.3.2 count_all()	236
8.40.3.3 count_current()	236
8.40.3.4 count_init()	236
8.40.3.5 get_counters()	236
8.40.3.6 get_descriptions()	236
8.40.3.7 get_names()	236
8.40.3.8 reset_array()	236
8.40.3.9 set_counters()	237
8.40.3.10 size()	237
8.41 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class Tem-	
	237
8.41.1 Detailed Description	
8.41.2 Constructor & Destructor Documentation	
8.41.2.1 Support() [1/3]	
<b>8.41.2.2 Support()</b> [2/3]	240
<b>8.41.2.3 Support()</b> [3/3]	240
8.41.2.4 ~Support()	240
8.41.3 Member Function Documentation	240
8.41.3.1 add_counter()	240
8.41.3.2 add_rule() [1/2]	241
8.41.3.3 add_rule() [2/2]	241
<b>8.41.3.4</b> add_rule_dyn() [1/2]	241
8.41.3.5 add_rule_dyn() [2/2]	241

8.41.3.6 calc()	2/1
8.41.3.7 eval rules dyn()	
8.41.3.8 get_counters()	
8.41.3.9 get_counts()	
8.41.3.10 get_current_stats()	
8.41.3.11 get data()	
8.41.3.12 get_rules()	
8.41.3.13 get_rules_dyn()	
8.41.3.14 init_support()	244
8.41.3.15 print()	244
8.41.3.16 reset_array() [1/2]	244
8.41.3.17 reset_array() [2/2]	244
8.41.3.18 set_counters()	245
8.41.3.19 set_rules()	245
8.41.3.20 set_rules_dyn()	245
8.41.4 Member Data Documentation	245
8.41.4.1 change_stats	245
8.41.4.2 coordiantes_n_free	246
8.41.4.3 coordiantes_n_locked	246
8.41.4.4 coordinates_free	246
8.41.4.5 coordinates_locked	246
8.41.4.6 current_stats	246
8.41.4.7 delete_counters	247
8.41.4.8 delete_rules	247
8.41.4.9 delete_rules_dyn	
8.41.4.10 hashes	247
8.41.4.11 hashes_initialized	247
8.41.4.12 M	248
8.41.4.13 max_num_elements	
8.41.4.14 N	
8.41.4.15 n_counters	
8.42 vecHasher< T > Struct Template Reference	
8.42.1 Detailed Description	
8.42.2 Member Function Documentation	
8.42.2.1 operator()()	249
9 File Documentation	251
9.1 include/barry/barray-bones.hpp File Reference	251
9.2 include/barry/barray-iterator.hpp File Reference	251
9.3 include/barry/barray-meat-operators.hpp File Reference	252
9.3.1 Macro Definition Documentation	252
9.3.1.1 BARRAY_TEMPLATE	253

9.3.1.2 BARRAY_TEMPLATE_ARGS
9.3.1.3 BARRAY_TYPE
9.3.1.4 COL
9.3.1.5 ROW
9.3.2 Function Documentation
9.3.2.1 BARRAY_TEMPLATE() [1/6]
9.3.2.2 BARRAY_TEMPLATE() [2/6]
9.3.2.3 BARRAY_TEMPLATE() [3/6]
9.3.2.4 BARRAY_TEMPLATE() [4/6]
9.3.2.5 BARRAY_TEMPLATE() [5/6]
9.3.2.6 BARRAY_TEMPLATE() [6/6]
9.3.2.7 BARRAY_TEMPLATE_ARGS()
9.3.2.8 BARRAY_TYPE()
9.3.2.9 for()
9.3.2.10 operator()()
9.3.3 Variable Documentation
9.3.3.1 rhs
9.3.3.2 this
9.4 include/barry/barray-meat.hpp File Reference
9.4.1 Macro Definition Documentation
9.4.1.1 COL
9.4.1.2 ROW
9.5 include/barry/barraycell-bones.hpp File Reference
9.6 include/barry/barraycell-meat.hpp File Reference
9.7 include/barry/barraydense-bones.hpp File Reference
9.8 include/barry/barraydense-meat-operators.hpp File Reference
9.8.1 Macro Definition Documentation
9.8.1.1 BDENSE_TEMPLATE
9.8.1.2 BDENSE_TEMPLATE_ARGS
9.8.1.3 BDENSE_TYPE
9.8.1.4 COL
9.8.1.5 POS
9.8.1.6 POS_N
9.8.1.7 ROW
9.8.2 Function Documentation
9.8.2.1 BDENSE_TEMPLATE() [1/4]
9.8.2.2 BDENSE_TEMPLATE() [2/4]
9.8.2.3 BDENSE_TEMPLATE() [3/4]
9.8.2.4 BDENSE_TEMPLATE() [4/4]
9.8.2.5 BDENSE_TEMPLATE_ARGS()
9.8.2.6 BDENSE_TYPE()
9.9 include/harry/harraydanse-meat hop File Reference

9.9.1 Macro Definition Documentation
9.9.1.1 COL
9.9.1.2 POS
9.9.1.3 POS_N
9.9.1.4 ROW
9.9.1.5 ZERO_CELL
9.10 include/barry/barraydensecell-bones.hpp File Reference
9.10.1 Macro Definition Documentation
9.10.1.1 POS
9.11 include/barry/barraydensecell-meat.hpp File Reference
9.11.1 Macro Definition Documentation
9.11.1.1 POS
9.12 include/barry/barraydensecol-bones.hpp File Reference
9.12.1 Macro Definition Documentation
9.12.1.1 POS
9.12.1.2 POS_N
9.12.1.3 ZERO_CELL
9.13 include/barry/barraydenserow-bones.hpp File Reference
9.13.1 Macro Definition Documentation
9.13.1.1 POS
9.13.1.2 POS_N
9.13.1.3 ZERO_CELL
9.14 include/barry/barrayrow-bones.hpp File Reference
9.15 include/barry/barrayrow-meat.hpp File Reference
9.15.1 Macro Definition Documentation
9.15.1.1 BROW_TEMPLATE
9.15.1.2 BROW_TEMPLATE_ARGS
9.15.1.3 BROW_TYPE
9.15.2 Function Documentation
9.15.2.1 BROW_TEMPLATE() [1/5]
9.15.2.2 BROW_TEMPLATE() [2/5]
9.15.2.3 BROW_TEMPLATE() [3/5]
9.15.2.4 BROW_TEMPLATE() [4/5]
9.15.2.5 BROW_TEMPLATE() [5/5]
9.16 include/barry/barrayvector-bones.hpp File Reference
9.17 include/barry/barrayvector-meat.hpp File Reference
9.18 include/barry/barry-configuration.hpp File Reference
9.18.1 Macro Definition Documentation
9.18.1.1 BARRY_CHECK_SUPPORT
9.18.1.2 BARRY_ISFINITE
9.18.1.3 BARRY_MAX_NUM_ELEMENTS
9.18.1.4 BARRY SAFE EXP

9.18.1.5 printf_barry
9.18.2 Typedef Documentation
9.18.2.1 Map
9.19 include/barry/barry-debug.hpp File Reference
9.19.1 Macro Definition Documentation
9.19.1.1 BARRY_DEBUG_LEVEL
9.20 include/barry/barry-macros.hpp File Reference
9.20.1 Macro Definition Documentation
9.20.1.1 BARRY_NCORES_ARG
9.20.1.2 BARRY_ONE
9.20.1.3 BARRY_ONE_DENSE
9.20.1.4 BARRY_UNUSED
9.20.1.5 BARRY_ZERO
9.20.1.6 BARRY_ZERO_DENSE
9.21 include/barry/barry.hpp File Reference
9.21.1 Macro Definition Documentation
9.21.1.1 BARRY_HPP
9.21.1.2 BARRY_VERSION
9.21.1.3 BARRY_VERSION_MAYOR
9.21.1.4 BARRY_VERSION_MINOR
9.21.1.5 COUNTER_FUNCTION
9.21.1.6 COUNTER_LAMBDA
9.21.1.7 RULE_FUNCTION
9.21.1.8 RULE_LAMBDA
9.22 include/barry/cell-bones.hpp File Reference
9.23 include/barry/cell-meat.hpp File Reference
9.24 include/barry/col-bones.hpp File Reference
9.25 include/barry/counters-bones.hpp File Reference
9.26 include/barry/counters-meat.hpp File Reference
9.26.1 Macro Definition Documentation
9.26.1.1 COUNTER_TEMPLATE
9.26.1.2 COUNTER_TEMPLATE_ARGS
9.26.1.3 COUNTER_TYPE
9.26.1.4 COUNTERS_TEMPLATE
9.26.1.5 COUNTERS_TEMPLATE_ARGS
9.26.1.6 COUNTERS_TYPE
9.26.1.7 TMP_HASHER_CALL
9.26.2 Function Documentation
9.26.2.1 count_fun()
9.26.2.2 COUNTER_TEMPLATE() [1/9]
9.26.2.3 COUNTER_TEMPLATE() [2/9]
9.26.2.4 COUNTER TEMPLATE() [3/9]

9.26.2.5 COUNTER_TEMPLATE() [4/9]	282
9.26.2.6 COUNTER_TEMPLATE() [5/9]	283
9.26.2.7 COUNTER_TEMPLATE() [6/9]	283
9.26.2.8 COUNTER_TEMPLATE() [7/9]	283
9.26.2.9 COUNTER_TEMPLATE() [8/9]	283
9.26.2.10 COUNTER_TEMPLATE() [9/9]	283
9.26.2.11 COUNTERS_TEMPLATE() [1/9]	284
9.26.2.12 COUNTERS_TEMPLATE() [2/9]	284
9.26.2.13 COUNTERS_TEMPLATE() [3/9]	284
9.26.2.14 COUNTERS_TEMPLATE() [4/9]	284
9.26.2.15 COUNTERS_TEMPLATE() [5/9]	284
9.26.2.16 COUNTERS_TEMPLATE() [6/9]	284
9.26.2.17 COUNTERS_TEMPLATE() [7/9]	285
9.26.2.18 COUNTERS_TEMPLATE() [8/9]	285
9.26.2.19 COUNTERS_TEMPLATE() [9/9]	285
9.26.2.20 data()	285
9.26.2.21 desc()	285
9.26.2.22 for()	285
9.26.2.23 hasher() [1/2]	286
9.26.2.24 hasher() [2/2]	286
9.26.2.25 hasher_fun() [1/2]	286
9.26.2.26 hasher_fun() [2/2]	286
9.26.2.27 if() [1/3]	286
9.26.2.28 if() [2/3]	286
9.26.2.29 if() [3/3]	287
9.26.2.30 init_fun() [1/3]	287
9.26.2.31 init_fun() [2/3]	287
9.26.2.32 init_fun() [3/3]	287
9.26.2.33 name()	287
9.26.3 Variable Documentation	287
9.26.3.1 add_dims	287
9.26.3.2 count_fun	288
9.26.3.3 counter	288
9.26.3.4 counter	288
9.26.3.5 data	288
9.26.3.6 desc	289
9.26.3.7 fun	289
9.26.3.8 fun	289
9.26.3.9 hasher_fun	289
9.26.3.10 i	290
9.26.3.11 init_fun	290
9.26.3.12 j	290

9.26.3.13 name	90
9.26.3.14 noexcept	90
9.26.3.15 res	91
9.26.3.16 return	91
9.27 include/barry/counters/network-css.hpp File Reference	91
9.27.1 Macro Definition Documentation	92
9.27.1.1 CSS_APPEND	93
9.27.1.2 CSS_CASE_ELSE	93
9.27.1.3 CSS_CASE_PERCEIVED	93
9.27.1.4 CSS_CASE_TRUTH	93
9.27.1.5 CSS_CHECK_SIZE	93
9.27.1.6 CSS_CHECK_SIZE_INIT	94
9.27.1.7 CSS_MATCH_TYPE	94
9.27.1.8 CSS_NET_COUNTER_LAMBDA_INIT	94
9.27.1.9 CSS_PERCEIVED_CELLS	94
9.27.1.10 CSS_SIZE	95
9.27.1.11 CSS_TRUE_CELLS	95
9.27.2 Function Documentation	95
9.27.2.1 counter_css_census01()	95
9.27.2.2 counter_css_census02()	95
9.27.2.3 counter_css_census03()	96
9.27.2.4 counter_css_census04()	96
9.27.2.5 counter_css_census05()	96
9.27.2.6 counter_css_census06()	96
9.27.2.7 counter_css_census07()	97
9.27.2.8 counter_css_census08()	97
9.27.2.9 counter_css_census09()	97
9.27.2.10 counter_css_census10()	97
9.27.2.11 counter_css_completely_false_recip_comiss()	98
9.27.2.12 counter_css_completely_false_recip_omiss()	98
9.27.2.13 counter_css_mixed_recip()	98
9.27.2.14 counter_css_partially_false_recip_commi()	98
9.27.2.15 counter_css_partially_false_recip_omiss()	99
9.28 include/barry/counters/network.hpp File Reference	99
9.28.1 Macro Definition Documentation	02
9.28.1.1 BARRY_ZERO_NETWORK	02
9.28.1.2 BARRY_ZERO_NETWORK_DENSE	03
9.28.1.3 NET_C_DATA_IDX	03
9.28.1.4 NET_C_DATA_NUM	03
9.28.1.5 NETWORK_COUNTER	03
9.28.1.6 NETWORK_COUNTER_LAMBDA	03
9.28.1.7 NETWORK_RULE	04

9.28.1.8 NETWORK_RULE_LAMBDA
9.28.1.9 NETWORKDENSE_COUNTER_LAMBDA
9.28.2 Typedef Documentation
9.28.2.1 NetCounter
9.28.2.2 NetCounters
9.28.2.3 NetModel
9.28.2.4 NetRule
9.28.2.5 NetRules
9.28.2.6 NetStatsCounter
9.28.2.7 NetSupport
9.28.2.8 Network
9.28.2.9 NetworkDense
9.28.3 Function Documentation
9.28.3.1 rules_zerodiag()
9.29 include/barry/freqtable.hpp File Reference
9.30 include/barry/model-bones.hpp File Reference
9.31 include/barry/model-meat.hpp File Reference
9.31.1 Function Documentation
9.31.1.1 likelihood_()
9.31.1.2 update_normalizing_constant()
9.32 include/barry/models/defm.hpp File Reference
9.33 include/barry/models/defm/counters.hpp File Reference
9.33.1 Macro Definition Documentation
9.33.1.1 DEFM_COUNTER
9.33.1.2 DEFM_COUNTER_LAMBDA
9.33.1.3 DEFM_RULE
9.33.1.4 DEFM_RULE_LAMBDA
9.33.1.5 DEFM_RULEDYN_LAMBDA
9.33.1.6 UNI_SUB
9.34 include/barry/models/geese/counters.hpp File Reference
9.35 include/barry/models/defm/defm-bones.hpp File Reference
9.36 include/barry/models/defm/defm-meat.hpp File Reference
9.36.1 Macro Definition Documentation
9.36.1.1 DEFM_LOOP_ARRAYS
9.36.1.2 DEFM_RANGES
9.36.2 Function Documentation
9.36.2.1 keygen_defm()
9.37 include/barry/models/defm/defm-types.hpp File Reference
9.37.1 Typedef Documentation
9.37.1.1 DEFMArray
9.38 include/barry/models/defm/formula.hpp File Reference
9.38.1 Function Documentation

9.38.1.1 defm_motif_parser()	18
9.39 include/barry/models/geese.hpp File Reference	19
9.40 include/barry/models/geese/flock-bones.hpp File Reference	19
9.41 include/barry/models/geese/flock-meat.hpp File Reference	20
9.42 include/barry/models/geese/geese-bones.hpp File Reference	20
9.42.1 Macro Definition Documentation	21
9.42.1.1 INITIALIZED	21
9.42.2 Function Documentation	21
9.42.2.1 keygen_full()	21
9.42.2.2 RULE_FUNCTION()	21
9.42.2.3 vec_diff()	22
9.42.2.4 vector_caster()	22
9.43 include/barry/models/geese/geese-meat-constructors.hpp File Reference	22
9.44 include/barry/models/geese/geese-meat-likelihood.hpp File Reference	23
9.44.1 Function Documentation	23
9.44.1.1 pset_loop()	24
9.45 include/barry/models/geese/geese-meat-likelihood_exhaust.hpp File Reference	24
9.46 include/barry/models/geese/geese-meat-predict.hpp File Reference	25
9.47 include/barry/models/geese/geese-meat-predict_exhaust.hpp File Reference	25
9.48 include/barry/models/geese/geese-meat-predict_sim.hpp File Reference	26
9.49 include/barry/models/geese/geese-meat-simulate.hpp File Reference	26
9.50 include/barry/models/geese/geese-meat.hpp File Reference	27
9.51 include/barry/models/geese/geese-node-bones.hpp File Reference	27
9.52 include/barry/models/geese/geese-types.hpp File Reference	28
9.52.1 Macro Definition Documentation	28
9.52.1.1 POS	29
9.52.2 Typedef Documentation	29
9.52.2.1 PhyloArray	29
9.52.2.2 PhyloCounter	29
9.52.2.3 PhyloCounters	29
9.52.2.4 PhyloModel	29
9.52.2.5 PhyloPowerSet	30
9.52.2.6 PhyloRule	30
9.52.2.7 PhyloRuleData	30
9.52.2.8 PhyloRuleDyn	30
9.52.2.9 PhyloRules	30
9.52.2.10 PhyloRulesDyn	30
9.52.2.11 PhyloStatsCounter	31
9.52.2.12 PhyloSupport	31
9.53 include/barry/powerset-bones.hpp File Reference	31
9.54 include/barry/powerset-meat.hpp File Reference	32
9.55 include/barry/progress.hpp File Reference	32

9.55.1 Macro Definition Documentation	3
9.55.1.1 BARRY_PROGRESS_BAR_WIDTH	3
9.56 include/barry/rules-bones.hpp File Reference	3
9.56.1 Function Documentation	3
9.56.1.1 rule_fun_default()	4
9.57 include/barry/rules-meat.hpp File Reference	4
9.58 include/barry/statscounter-bones.hpp File Reference	4
9.59 include/barry/statscounter-meat.hpp File Reference	5
9.59.1 Macro Definition Documentation	6
9.59.1.1 STATSCOUNTER_TEMPLATE	6
9.59.1.2 STATSCOUNTER_TEMPLATE_ARGS	6
9.59.1.3 STATSCOUNTER_TYPE	6
9.59.2 Function Documentation	6
9.59.2.1 clear()	6
9.59.2.2 for()	7
9.59.2.3 resize()	7
9.59.2.4 STATSCOUNTER_TEMPLATE() [1/9]	7
9.59.2.5 STATSCOUNTER_TEMPLATE() [2/9]	7
9.59.2.6 STATSCOUNTER_TEMPLATE() [3/9]	7
9.59.2.7 STATSCOUNTER_TEMPLATE() [4/9]	7
9.59.2.8 STATSCOUNTER_TEMPLATE() [5/9]	8
9.59.2.9 STATSCOUNTER_TEMPLATE() [6/9]	8
9.59.2.10 STATSCOUNTER_TEMPLATE() [7/9]	8
9.59.2.11 STATSCOUNTER_TEMPLATE() [8/9]	8
9.59.2.12 STATSCOUNTER_TEMPLATE() [9/9]	8
9.59.3 Variable Documentation	8
9.59.3.1 counter	9
9.59.3.2 counter_deleted	9
9.59.3.3 counters	9
9.59.3.4 counters	9
9.59.3.5 current_stats	9
9.59.3.6 EmptyArray	0
9.59.3.7 f	0
9.59.3.8 j	0
9.59.3.9 return	0
9.60 include/barry/support-bones.hpp File Reference	0
9.61 include/barry/support-meat.hpp File Reference	-1
9.61.1 Macro Definition Documentation	-1
9.61.1.1 BARRY_SUPPORT_MEAT_HPP	.1
9.62 include/barry/typedefs.hpp File Reference	2
9.62.1 Typedef Documentation	4
9.62.1.1 Col. type	4

9.62.1.2 Counter_fun_type	344
9.62.1.3 Counts_type	344
9.62.1.4 Hasher_fun_type	344
9.62.1.5 MapVec_type	345
9.62.1.6 Row_type	345
9.62.1.7 Rule_fun_type	345
9.62.2 Function Documentation	345
9.62.2.1 sort_array()	345
9.62.2.2 vec_equal()	346
9.62.2.3 vec_equal_approx()	346
9.62.2.4 vec_inner_prod() [1/2]	347
9.62.2.5 vec_inner_prod() [2/2]	347
9.63 README.md File Reference	347
Index	349

# **Chapter 1**

# Main Page

## Barry: your to-go motif accountant

This repository contains a C++ template library that essentially counts sufficient statistics on binary arrays. Its primary goal is to provide a general framework for building discrete exponential-family models. A particular example is Exponential Random Graph Models (ERGMs), but we can use barry to deal with non-square arrays.

Among the key features included in barry, we have:

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

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

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

2 Main Page

## **Examples**

#### Counting statistics in a graph

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

```
#include <ostream>
#include "../include/barry.hpp"
typedef std::vector< unsigned int > vuint;
int main() {
  // Creating network of size six with five ties
  netcounters::Network net(
       {0, 0, 4, 4, 2, 0, 1},
       {1, 2, 0, 2, 4, 0, 1}
  // How does this looks like?
  net.print("Current view");
  // Adding extra ties
  net += \{1, 0\};
  net(2, 0) = true;
  // And removing a couple
  net(0, 0) = false;
net -= {1, 1};
net.print("New view");
  // Initializing the data. The program deals with freing the memory
  net.set_data(new netcounters::NetworkData, true);
  // Creating counter object for the network and adding stats to count
  netcounters::NetStatsCounter counter(&net);
  netcounters::counter_edges(counter.counters);
  netcounters::counter_ttriads(counter.counters);
  netcounters::counter_isolates(counter.counters);
  netcounters::counter_ctriads(counter.counters);
  netcounters::counter_mutual(counter.counters);
  // Counting and printing the results
std::vector< double > counts = counter.count_all();
    "Edges : " « counts[0] « std::endl «
"Transitive triads : " « counts[1] « std::endl «
"Isolates : " « counts[2] « std::endl «
"C triads : " « counts[3] « std::endl «
"Mutuals : " « counts[4] « std::endl;
  return 0;
```

#### Compiling this program using g++

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

#### Yields the following output:

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

### **Features**

### **Efficient memory usage**

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

### **Documentation**

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

#### **Code of Conduct**

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

4 Main Page

# **Chapter 2**

# **Module Index**

# 2.1 Modules

Here is a list of all modules:

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

6 Module Index

# **Chapter 3**

# **Hierarchical Index**

# 3.1 Class Hierarchy

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

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

8 Hierarchical Index

# **Chapter 4**

# **Class Index**

## 4.1 Class List

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

BArray < Cell_Type, Data_Type >
Baseline class for binary arrays
BArrayCell < Cell_Type, Data_Type >
BArrayCell_const< Cell_Type, Data_Type >
BArrayDense < Cell_Type, Data_Type >
Baseline class for binary arrays
BArrayDenseCell< Cell_Type, Data_Type >
BArrayDenseCell_const< Cell_Type, Data_Type >
BArrayDenseCol < Cell_Type, Data_Type >
BArrayDenseCol_const< Cell_Type, Data_Type >
BArrayDenseRow< Cell_Type, Data_Type >
BArrayDenseRow_const< Cell_Type, Data_Type >
BArrayRow< Cell_Type, Data_Type >
BArrayRow_const< Cell_Type, Data_Type >
BArrayVector< Cell_Type, Data_Type >
Row or column of a BArray 113
BArrayVector_const< Cell_Type, Data_Type >
Cell< Cell_Type >
Entries in BArray. For now, it only has two members:
Cell_const< Cell_Type >
ConstBArrayRowlter < Cell_Type, Data_Type >
Counter< Array_Type, Data_Type >
A counter function based on change statistics
Counters < Array_Type, Data_Type >
Vector of counters
DEFM 139
DEFMCounterData
Data class used to store arbitrary size_t or double vectors
DEFMData
Data class for DEFM arrays
DEFMRuleData
DEFMRuleDynData
Entries < Cell_Type >
A wrapper class to store source, target, val from a BArray object
Flock
A Flock is a group of Geese

10 Class Index

FreqTable< T >	
Frequency table of vectors	157
Geese	
Annotated Phylo Model	160
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:	175
NetCounterData	
Data class used to store arbitrary size_t or double vectors	202
NetworkData	
Data class for Networks	203
Node	
A single node for the model	206
NodeData	
Data definition for the PhyloArray class	212
PhyloCounterData	213
PhyloRuleDynData	216
PowerSet< Array_Type, Data_Rule_Type >	
Powerset of a binary array	219
Progress	
A simple progress bar	225
Rule < Array_Type, Data_Type >	
Rule for determining if a cell should be included in a sequence	226
Rules< Array_Type, Data_Type >	
Vector of objects of class Rule	229
StatsCounter< Array_Type, Data_Type >	
Count stats for a single Array	233
Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >	
Compute the support of sufficient statistics	237
vecHasher <t></t>	248

# **Chapter 5**

# File Index

# 5.1 File List

Here is a list of all files with brief descriptions:

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

12 File Index

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

# **Chapter 6**

# **Module Documentation**

## 6.1 Counting

#### **Classes**

· class NetworkData

Data class for Networks.

class Counter< Array\_Type, Data\_Type >

A counter function based on change statistics.

#### **Macros**

- #define MAKE\_DEFM\_HASHER(hasher, a, cov)
- #define MAKE\_DUPL\_VARS()
- #define IS\_EITHER() (DATA\_AT == Geese::etype\_either)
- #define IS\_DUPLICATION() ((DATA\_AT == Geese::etype\_duplication) & (DPL))
- #define IS\_SPECIATION() ((DATA\_AT == Geese::etype\_speciation) & (!DPL))
- #define IF\_MATCHES()
- #define IF NOTMATCHES()
- #define PHYLO RULE LAMBDA(a)

Extension of a simple counter.

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

Overall functional gains.

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

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

k genes gain function nfun

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

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

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

Keeps track of how many pairs of genes preserve pseudostate.

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

Overall functional loss.

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

6.1 Counting 15

#### 6.1.1 Detailed Description

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

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

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

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

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

#### 6.1.2 Macro Definition Documentation

#### **6.1.2.1 IF MATCHES**

Definition at line 20 of file counters.hpp.

#### 6.1.2.2 IF\_NOTMATCHES

```
#define IF_NOTMATCHES( )

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

Definition at line 22 of file counters.hpp.

#### 6.1.2.3 IS\_DUPLICATION

Definition at line 17 of file counters.hpp.

#### 6.1.2.4 IS\_EITHER

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

Definition at line 16 of file counters.hpp.

#### 6.1.2.5 IS\_SPECIATION

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

Definition at line 18 of file counters.hpp.

#### 6.1.2.6 MAKE DEFM HASHER

#### Value:

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

Definition at line 21 of file counters.hpp.

#### 6.1.2.7 MAKE\_DUPL\_VARS

```
#define MAKE_DUPL_VARS( )

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

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

Definition at line 12 of file counters.hpp.

6.1 Counting 17

#### 6.1.2.8 PHYLO\_CHECK\_MISSING

```
#define PHYLO_CHECK_MISSING( )

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

Definition at line 45 of file counters.hpp.

#### 6.1.2.9 PHYLO\_COUNTER\_LAMBDA

Definition at line 39 of file counters.hpp.

#### 6.1.2.10 PHYLO\_RULE\_DYN\_LAMBDA

Definition at line 42 of file counters.hpp.

#### 6.1.2.11 PHYLO RULE LAMBDA

Extension of a simple counter.

It allows specifying extra arguments, in particular, the corresponding sets of rows to which this statistic may be relevant. This could be important in the case of, for example, counting correlation type statistics between function 1 and 2, and between function 1 and 3.

Definition at line 36 of file counters.hpp.

#### 6.1.3 Function Documentation

#### 6.1.3.1 counter\_co\_opt()

Function co-opting.

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

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

This algorithm implements the change statistic.

Definition at line 1299 of file counters.hpp.

#### 6.1.3.2 counter\_cogain()

Co-evolution (joint gain or loss)

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

Definition at line 794 of file counters.hpp.

### 6.1.3.3 counter\_gains()

Functional gains for a specific function (nfun).

Definition at line 99 of file counters.hpp.

6.1 Counting 19

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

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

Needs to specify function a.

Definition at line 1633 of file counters.hpp.

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

k genes gain function nfun

Definition at line 159 of file counters.hpp.

#### 6.1.3.6 counter\_genes\_changing()

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

Definition at line 231 of file counters.hpp.

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

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

Definition at line 1397 of file counters.hpp.

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

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

< How many genes diverge the parent

Definition at line 1517 of file counters.hpp.

#### 6.1.3.9 counter\_longest()

Longest branch mutates (either by gain or by loss)

Definition at line 851 of file counters.hpp.

#### 6.1.3.10 counter\_loss()

Total count of losses for an specific function.

Definition at line 594 of file counters.hpp.

#### 6.1.3.11 counter\_maxfuns()

Cap the number of functions per gene.

Definition at line 532 of file counters.hpp.

6.1 Counting 21

#### 6.1.3.12 counter\_neofun()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1021 of file counters.hpp.

#### 6.1.3.13 counter\_neofun\_a2b()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1166 of file counters.hpp.

#### 6.1.3.14 counter\_overall\_changes()

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

Definition at line 646 of file counters.hpp.

#### 6.1.3.15 counter\_overall\_gains()

Overall functional gains.

Total number of gains (irrespective of the function).

Definition at line 61 of file counters.hpp.

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

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

Needs to specify function a.

Definition at line 1699 of file counters.hpp.

#### 6.1.3.17 counter\_overall\_loss()

Overall functional loss.

Definition at line 484 of file counters.hpp.

#### 6.1.3.18 counter pairwise first gain()

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

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

Definition at line 1951 of file counters.hpp.

#### 6.1.3.19 counter pairwise neofun singlefun()

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

Definition at line 1102 of file counters.hpp.

6.1 Counting 23

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

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

Needs to specify function a.

Definition at line 1747 of file counters.hpp.

#### 6.1.3.21 counter pairwise preserving()

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

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

Definition at line 1812 of file counters.hpp.

#### 6.1.3.22 counter\_preserve\_pseudogene()

Keeps track of how many pairs of genes preserve pseudostate.

Definition at line 300 of file counters.hpp.

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

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

Definition at line 382 of file counters.hpp.

#### 6.1.3.24 counter\_subfun()

Total count of Sub-functionalization events.

It requires to specify data = {funA, funB}

Definition at line 705 of file counters.hpp.

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

Definition at line 48 of file counters.hpp.

#### 6.2 Statistical Models

Statistical models available in barry.

#### Classes

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

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

class Flock

A Flock is a group of Geese.

· class Geese

Annotated Phylo Model.

#### 6.2.1 Detailed Description

Statistical models available in barry.

### 6.3 Network counters

Counters for network models.

6.3 Network counters 25

#### **Functions**

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

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

    template<typename Tnet = Network>

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

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

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

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

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

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

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

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

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

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

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

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

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

    NETWORK_COUNTER (init_single_attr)

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

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

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

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

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

Counts number of vertices with a given out-degree.

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

Prevalence of ones.

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

Prevalence of ones.

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

Prevalence of ones.

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

Prevalence of ones.

#### Returns true if the cell is free

#### **Parameters**

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

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

Number of edges.

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

Blocks switching a one to zero.

#### 6.3.1 Detailed Description

Counters for network models.

#### **Parameters**

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

#### 6.3.2 Function Documentation

6.3 Network counters 27

#### 6.3.2.1 counter\_absdiff()

Sum of absolute attribute difference between ego and alter.

Definition at line 908 of file network.hpp.

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

Definition at line 665 of file network.hpp.

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

Definition at line 610 of file network.hpp.

#### 6.3.2.4 counter\_degree()

Counts number of vertices with a given out-degree.

Definition at line 1326 of file network.hpp.

#### 6.3.2.5 counter\_density()

Definition at line 729 of file network.hpp.

#### 6.3.2.6 counter\_diff()

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

Definition at line 953 of file network.hpp.

#### 6.3.2.7 counter\_edges()

Number of edges.

Definition at line 152 of file network.hpp.

#### 6.3.2.8 counter\_fixed\_effect()

Prevalence of ones.

#### **Parameters**

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

6.3 Network counters 29

Definition at line 610 of file counters.hpp.

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

Definition at line 1170 of file network.hpp.

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

Counts number of vertices with a given in-degree.

Definition at line 1123 of file network.hpp.

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

Definition at line 785 of file network.hpp.

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

Definition at line 757 of file network.hpp.

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

Definition at line 215 of file network.hpp.

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

Number of isolated vertices.

Definition at line 175 of file network.hpp.

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

Definition at line 338 of file network.hpp.

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

Definition at line 312 of file network.hpp.

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

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

Definition at line 151 of file counters.hpp.

6.3 Network counters 31

#### 6.3.2.18 counter\_mutual()

Number of mutual ties.

Definition at line 256 of file network.hpp.

#### 6.3.2.19 counter\_nodecov()

Definition at line 1066 of file network.hpp.

#### 6.3.2.20 counter\_nodeicov()

Definition at line 1016 of file network.hpp.

#### 6.3.2.21 counter\_nodematch()

Definition at line 1091 of file network.hpp.

#### 6.3.2.22 counter\_nodeocov()

Definition at line 1041 of file network.hpp.

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

Definition at line 1271 of file network.hpp.

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

Counts number of vertices with a given out-degree.

Definition at line 1223 of file network.hpp.

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

Definition at line 862 of file network.hpp.

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

Definition at line 834 of file network.hpp.

#### 6.3.2.27 counter\_ones()

Prevalence of ones.

6.3 Network counters 33

#### **Parameters**

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

Definition at line 81 of file counters.hpp.

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

Definition at line 404 of file network.hpp.

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

Definition at line 376 of file network.hpp.

#### 6.3.2.30 counter\_transition()

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

Prevalence of ones.

#### **Parameters**

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

Definition at line 270 of file counters.hpp.

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

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

Prevalence of ones.

#### **Parameters**

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

Definition at line 579 of file counters.hpp.

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

Definition at line 531 of file network.hpp.

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

Definition at line 441 of file network.hpp.

#### 6.3.2.34 NETWORK\_COUNTER()

Definition at line 997 of file network.hpp.

6.4 Phylo rules 35

#### 6.3.2.35 rules\_dont\_become\_zero()

Blocks switching a one to zero.

#### **Parameters**

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

Definition at line 678 of file counters.hpp.

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

Number of edges.

Definition at line 653 of file counters.hpp.

## 6.4 Phylo rules

Rules for phylogenetic modeling.

Collaboration diagram for Phylo rules:



#### Classes

• class DEFMRuleDynData

#### **Functions**

DEFMData::DEFMData ()

Vector indicating which covariates are included in the model.

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

Constructor.

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

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

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

Overall functional gains.

## **Variables**

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

Vector of covariates (complete vector)

· size t DEFMData::obs start

Index of the observation in the data.

size\_t DEFMData::X\_ncol

Number of columns in the array of covariates.

size t DEFMData::X nrow

Number of rows in the array of covariates.

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

Value where the sorting of the covariates is stored.

- std::vector< size t > DEFMCounterData::indices
- std::vector< double > DEFMCounterData::numbers
- std::vector< bool > DEFMCounterData::logical

6.4 Phylo rules 37

- · bool DEFMCounterData::is\_motif
  - If false, then is a logit intercept.
- std::vector< double > DEFMRuleData::numbers
- std::vector< size t > DEFMRuleData::indices
- std::vector< bool > DEFMRuleData::logical
- bool DEFMRuleData::init = false
- const std::vector< double > \* DEFMRuleDynData::counts

## Convenient typedefs for network objects.

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

## 6.4.1 Detailed Description

Rules for phylogenetic modeling.

#### **Parameters**

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

## 6.4.2 Typedef Documentation

#### 6.4.2.1 DEFMCounter

typedef barry::Counter<DEFMArray, DEFMCounterData > DEFMCounter

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

## 6.4.2.2 DEFMCounters

typedef barry::Counters<DEFMArray, DEFMCounterData> DEFMCounters

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

## 6.4.2.3 DEFMModel

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

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

#### 6.4.2.4 DEFMRule

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

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

## 6.4.2.5 DEFMRuleDyn

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

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

## 6.4.2.6 DEFMRules

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

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

## 6.4.2.7 DEFMRulesDyn

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

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

#### 6.4.2.8 DEFMStatsCounter

typedef barry::StatsCounter<DEFMArray, DEFMCounterData> DEFMStatsCounter

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

6.4 Phylo rules 39

#### 6.4.2.9 DEFMSupport

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

#### 6.4.3 Function Documentation

## 6.4.3.1 at()

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

## 6.4.3.2 DEFMCounterData() [1/2]

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

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

## 6.4.3.3 DEFMCounterData() [2/2]

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

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

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

Vector indicating which covariates are included in the model.

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

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

Constructor.

#### **Parameters**

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

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

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

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

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

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

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

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

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

## 6.4.3.9 DEFMRuleDynData()

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

6.4 Phylo rules 41

## 6.4.3.10 idx() [1/2]

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

## 6.4.3.11 idx() [2/2]

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

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

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

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

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

## 6.4.3.14 ncol()

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

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

## 6.4.3.15 nrow()

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

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

## 6.4.3.16 num() [1/2]

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

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

## 6.4.3.17 num() [2/2]

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

## 6.4.3.18 operator()()

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

## **Parameters**



Returns

double

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

## 6.4.3.19 print()

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

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

6.4 Phylo rules 43

## 6.4.3.20 rule\_dyn\_limit\_changes()

Overall functional gains.

#### **Parameters**

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

#### Returns

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

Definition at line 2087 of file counters.hpp.

## 6.4.3.21 rule\_leafs()

Definition at line 2052 of file counters.hpp.

## 6.4.3.22 ∼DEFMCounterData()

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

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

## 6.4.3.23 ∼DEFMData()

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

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

## 6.4.3.24 ∼DEFMRuleDynData()

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

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

## 6.4.4 Variable Documentation

## 6.4.4.1 array

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

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

## 6.4.4.2 counts

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

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

## 6.4.4.3 covar\_sort

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

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

## 6.4.4.4 covar\_used

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

Value where the sorting of the covariates is stored.

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

6.4 Phylo rules 45

## 6.4.4.5 covariates

const double\* DEFMData::covariates

Vector of covariates (complete vector)

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

## 6.4.4.6 indices [1/2]

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

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

#### 6.4.4.7 indices [2/2]

std::vector< size\_t > DEFMRuleData::indices

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

#### 6.4.4.8 init

bool DEFMRuleData::init = false

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

## 6.4.4.9 is\_motif

bool DEFMCounterData::is\_motif

If false, then is a logit intercept.

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

## 6.4.4.10 logical [1/2]

std::vector< bool > DEFMCounterData::logical

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

## 6.4.4.11 logical [2/2]

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

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

## 6.4.4.12 numbers [1/2]

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

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

#### 6.4.4.13 numbers [2/2]

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

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

## 6.4.4.14 obs\_start

```
size_t DEFMData::obs_start
```

Index of the observation in the data.

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

## 6.4.4.15 X\_ncol

```
size_t DEFMData::X_ncol
```

Number of columns in the array of covariates.

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

## 6.4.4.16 X\_nrow

```
size_t DEFMData::X_nrow
```

Number of rows in the array of covariates.

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

6.5 Phylo counters 47

## 6.5 Phylo counters

Counters for phylogenetic modeling.

Collaboration diagram for Phylo counters:



#### **Modules**

· Phylo rules

Rules for phylogenetic modeling.

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

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

k genes gain function nfun

- $\bullet \ \ void\ counter\_genes\_changing\ (PhyloCounters *counters, size\_t\ duplication=Geese::etype\_default)$
- void counter\_preserve\_pseudogene (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_ t duplication=Geese::etype default)

Keeps track of how many pairs of genes preserve pseudostate.

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

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

- void counter\_overall\_loss (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default) Overall functional loss.
- void counter\_maxfuns (PhyloCounters \*counters, size\_t lb, size\_t ub, size\_t duplication=Geese::etype\_default)

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

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

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

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

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

Total number of neofunctionalization events.

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

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

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

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

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

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

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

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

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

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

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

## 6.5.1 Detailed Description

Counters for phylogenetic modeling.

**Parameters** 

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

## 6.5.2 Function Documentation

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

Function co-opting.

6.5 Phylo counters 49

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

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

This algorithm implements the change statistic.

Definition at line 1299 of file counters.hpp.

#### 6.5.2.2 counter\_cogain()

Co-evolution (joint gain or loss)

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

Definition at line 794 of file counters.hpp.

## 6.5.2.3 counter\_gains()

Functional gains for a specific function (nfun).

Definition at line 99 of file counters.hpp.

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

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

Needs to specify function a.

Definition at line 1633 of file counters.hpp.

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

k genes gain function nfun

Definition at line 159 of file counters.hpp.

## 6.5.2.6 counter\_genes\_changing()

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

Definition at line 231 of file counters.hpp.

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

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

Definition at line 1397 of file counters.hpp.

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

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

< How many genes diverge the parent

Definition at line 1517 of file counters.hpp.

6.5 Phylo counters 51

#### 6.5.2.9 counter\_longest()

Longest branch mutates (either by gain or by loss)

Definition at line 851 of file counters.hpp.

## 6.5.2.10 counter\_loss()

Total count of losses for an specific function.

Definition at line 594 of file counters.hpp.

#### 6.5.2.11 counter\_maxfuns()

Cap the number of functions per gene.

Definition at line 532 of file counters.hpp.

## 6.5.2.12 counter\_neofun()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1021 of file counters.hpp.

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

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1166 of file counters.hpp.

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

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

Definition at line 646 of file counters.hpp.

## 6.5.2.15 counter\_overall\_gains()

Overall functional gains.

Total number of gains (irrespective of the function).

Definition at line 61 of file counters.hpp.

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

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

Needs to specify function a.

Definition at line 1699 of file counters.hpp.

6.5 Phylo counters 53

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

Overall functional loss.

Definition at line 484 of file counters.hpp.

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

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

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

Definition at line 1951 of file counters.hpp.

## 6.5.2.19 counter\_pairwise\_neofun\_singlefun()

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

Definition at line 1102 of file counters.hpp.

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

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

Needs to specify function a.

Definition at line 1747 of file counters.hpp.

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

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

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

Definition at line 1812 of file counters.hpp.

## 6.5.2.22 counter\_preserve\_pseudogene()

Keeps track of how many pairs of genes preserve pseudostate.

Definition at line 300 of file counters.hpp.

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

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

Definition at line 382 of file counters.hpp.

## 6.5.2.24 counter\_subfun()

Total count of Sub-functionalization events.

It requires to specify data = {funA, funB}

Definition at line 705 of file counters.hpp.

# **Chapter 7**

# **Namespace Documentation**

## 7.1 barry Namespace Reference

barry: Your go-to motif accountant

## **Namespaces**

counters

Tree class and Treelterator class.

## 7.1.1 Detailed Description

barry: Your go-to motif accountant

## 7.2 barry::counters Namespace Reference

Tree class and Treelterator class.

## **Namespaces**

network

## 7.2.1 Detailed Description

Tree class and Treelterator class.

## 7.3 barry::counters::network Namespace Reference

## 7.4 CHECK Namespace Reference

Integer constants used to specify which cell should be check.

## **Variables**

- const int BOTH = -1
  const int NONE = 0
  const int ONE = 1
  const int TWO = 2
- 7.4.1 Detailed Description

Integer constants used to specify which cell should be check.

## 7.4.2 Variable Documentation

#### 7.4.2.1 BOTH

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

Definition at line 27 of file typedefs.hpp.

## 7.4.2.2 NONE

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

Definition at line 28 of file typedefs.hpp.

### 7.4.2.3 ONE

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

Definition at line 29 of file typedefs.hpp.

## 7.4.2.4 TWO

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

Definition at line 30 of file typedefs.hpp.

## 7.5 defm Namespace Reference

## 7.6 EXISTS Namespace Reference

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

#### **Variables**

```
• const int BOTH = -1
```

- const int NONE = 0
- const int ONE = 1
- const int TWO = 1
- const int UKNOWN = -1
- const int AS\_ZERO = 0
- const int AS\_ONE = 1

## 7.6.1 Detailed Description

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

## 7.6.2 Variable Documentation

## 7.6.2.1 AS\_ONE

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

Definition at line 45 of file typedefs.hpp.

## 7.6.2.2 AS\_ZERO

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

Definition at line 44 of file typedefs.hpp.

## 7.6.2.3 BOTH

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

Definition at line 38 of file typedefs.hpp.

#### 7.6.2.4 NONE

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

Definition at line 39 of file typedefs.hpp.

#### 7.6.2.5 ONE

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

Definition at line 40 of file typedefs.hpp.

## 7.6.2.6 TWO

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

Definition at line 41 of file typedefs.hpp.

## 7.6.2.7 UKNOWN

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

Definition at line 43 of file typedefs.hpp.

## 7.7 geese Namespace Reference

## **Chapter 8**

## **Class Documentation**

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

Baseline class for binary arrays.

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

#### **Public Member Functions**

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

## Get the edgelist.

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

#### Constructors

#### **Parameters**

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

60 Class Documentation

• BArray ()

Zero-size array.

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

Empty array.

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

Edgelist with data.

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

Edgelist with no data (simpler)

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

Move operator.

- BArray< Cell\_Type, Data\_Type > & operator= (BArray< Cell\_Type, Data\_Type > &&x) noexcept
   Move assignment.
- void set\_data (Data\_Type \*data\_, bool delete\_data\_=false)

Set the data object.

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

## Queries

is\_empty queries a single cell. nrow, ncol, and nnozero return the number of rows, columns, and non-zero cells respectively.

#### **Parameters**

i,j	Coordinates
check_bounds	If false avoids checking bounds.

- bool is\_empty (size\_t i, size\_t j, bool check\_bounds=true) const
- size\_t nrow () const noexcept
- size\_t ncol () const noexcept
- size\_t nnozero () const noexcept
- Cell< Cell\_Type > default\_val () const

## Cell-wise insertion/deletion

## Parameters

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

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

## Column/row wise interchange

- void swap rows (size t i0, size t i1, bool check bounds=true)
- void swap\_cols (size\_t j0, size\_t j1, bool check bounds=true)
- void zero row (size t i, bool check bounds=true)
- void zero col (size t j, bool check bounds=true)

## **Arithmetic operators**

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

#### **Public Attributes**

· bool visited = false

#### **Friends**

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

#### 8.1.1 Detailed Description

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

Baseline class for binary arrays.

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

#### **Template Parameters**

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

62 Class Documentation

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

## 8.1.2 Constructor & Destructor Documentation

## 8.1.2.1 BArray() [1/6]

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

Zero-size array.

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

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

Empty array.

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

## 8.1.2.3 BArray() [3/6]

Edgelist with data.

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

Edgelist with no data (simpler)

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

Copy constructor.

## 8.1.2.6 BArray() [6/6]

Move operator.

## 8.1.2.7 $\sim$ BArray()

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

## 8.1.3 Member Function Documentation

## 8.1.3.1 clear()

64 Class Documentation

## 8.1.3.2 col()

## 8.1.3.3 D() [1/2]

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

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

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

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

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

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

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

## 8.1.3.7 default\_val()

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

## 8.1.3.8 flush\_data()

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

#### 8.1.3.9 get\_cell()

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

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

## 8.1.3.12 get\_entries()

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

## Get the edgelist.

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

#### Returns

Entries < Cell\_Type >

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

66 Class Documentation

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

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

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

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

#### 8.1.3.18 is dense()

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

Definition at line 241 of file barray-bones.hpp.

#### 8.1.3.19 is\_empty()

## 8.1.3.20 ncol()

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

#### 8.1.3.21 nnozero()

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

## 8.1.3.22 nrow()

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

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

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

68 Class Documentation

#### 8.1.3.25 operator\*=()

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

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

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

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

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

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

## 8.1.3.32 operator/=()

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

Move assignment.

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

Assignment constructor.

#### 8.1.3.35 operator==()

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

70 Class Documentation

## 8.1.3.37 print()

#### 8.1.3.38 print\_n()

#### 8.1.3.39 reserve()

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

## 8.1.3.40 resize()

#### 8.1.3.41 rm\_cell()

#### 8.1.3.42 row()

## 8.1.3.43 set\_data()

Set the data object.

#### **Parameters**

data_	
delete_ <i>←</i>	
data_	

### 8.1.3.44 swap\_cells()

## 8.1.3.45 swap\_cols()

72 Class Documentation

#### 8.1.3.46 swap\_rows()

## 8.1.3.47 toggle\_cell()

#### 8.1.3.48 toggle\_lock()

## 8.1.3.49 transpose()

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

## 8.1.3.50 zero\_col()

#### 8.1.3.51 zero\_row()

#### 8.1.4 Friends And Related Function Documentation

## 8.1.4.1 BArrayCell< Cell\_Type, Data\_Type >

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

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

## 8.1.4.2 BArrayCell\_const < Cell\_Type, Data\_Type >

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

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

## 8.1.5 Member Data Documentation

### 8.1.5.1 visited

```
template<typename Cell_Type = bool, typename Data_Type = bool>
bool BArray< Cell_Type, Data_Type >::visited = false
```

This is as a reference, if we need to iterate through the cells and we need to keep track which were visited, we use this as a reference. So that if cell.visited = true and visited = true, it means that we haven't been here yet. Ideally, any routine using this->visited should switch it at the beginning of the routine.

Definition at line 54 of file barray-bones.hpp.

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

• include/barry/barray-bones.hpp

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

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

#### **Public Member Functions**

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

# 8.2.1 Detailed Description

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

• bool operator== (const Cell\_Type &val) const

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

## 8.2.2 Constructor & Destructor Documentation

### 8.2.2.1 BArrayCell()

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

## 8.2.2.2 ~BArrayCell()

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

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

## 8.2.3 Member Function Documentation

## 8.2.3.1 operator Cell\_Type()

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

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

#### 8.2.3.2 operator\*=()

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

#### 8.2.3.3 operator+=()

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

#### 8.2.3.4 operator-=()

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

#### 8.2.3.5 operator/=()

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

#### 8.2.3.6 operator=()

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

#### 8.2.3.7 operator==()

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

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

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

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

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

## **Public Member Functions**

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

## 8.3.1 Detailed Description

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

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

## 8.3.2 Constructor & Destructor Documentation

## 8.3.2.1 BArrayCell\_const()

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

#### 8.3.2.2 ~BArrayCell\_const()

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

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

#### 8.3.3 Member Function Documentation

#### 8.3.3.1 operator Cell\_Type()

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

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

#### 8.3.3.2 operator"!=()

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

## 8.3.3.3 operator<()

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

#### 8.3.3.4 operator<=()

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

#### 8.3.3.5 operator==()

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

#### 8.3.3.6 operator>()

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

# 8.3.3.7 operator>=()

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

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

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

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

Baseline class for binary arrays.

#include <barraydense-bones.hpp>

#### **Public Member Functions**

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

Get the edgelist.

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

#### Constructors

#### **Parameters**

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

• BArrayDense ()

Zero-size array.

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

Edgelist with no data (simpler)

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

Assignment constructor.

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

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

Move assignment.

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

Set the data object.

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

#### Queries

is\_empty queries a single cell. nrow, ncol, and nnozero return the number of rows, columns, and non-zero cells respectively.

#### **Parameters**

i,j	Coordinates
check_bounds	If false avoids checking bounds.

- bool is\_empty (size\_t i, size\_t j, bool check\_bounds=true) const
- size t nrow () const noexcept
- size\_t ncol () const noexcept
- size\_t nnozero () const noexcept
- Cell< Cell\_Type > default\_val () const

### Cell-wise insertion/deletion

#### Parameters

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

- BArrayDense< Cell\_Type, Data\_Type > & operator+= (const std::pair< size\_t, size\_t > &coords)
- $\bullet \ \, \mathsf{BArrayDense} < \mathsf{Cell\_Type}, \, \mathsf{Data\_Type} > \& \, \mathsf{operator}\text{--=} \, (\mathsf{const} \, \, \mathsf{std} \text{::pair} < \, \mathsf{size\_t}, \, \mathsf{size\_t} > \& \mathsf{coords})$
- BArrayDenseCell< Cell\_Type, Data\_Type > operator() (size\_t i, size\_t j, bool check\_bounds=true)
- const Cell\_Type operator() (size\_t i, size\_t j, bool check\_bounds=true) const
- void rm\_cell (size\_t i, size\_t j, bool check\_bounds=true, bool check\_exists=true)
- void insert\_cell (size t i, size t j, const Cell< Cell Type > &v, bool check\_bounds, bool)
- void insert cell (size t i, size t j, Cell Type v, bool check bounds, bool)

- void swap\_cells (size\_t i0, size\_t j0, size\_t i1, size\_t j1, bool check\_bounds=true, int check\_
   exists=CHECK::BOTH, int \*report=nullptr)
- void toggle cell (size t i, size t j, bool check bounds=true, int check exists=EXISTS::UKNOWN)
- void toggle\_lock (size\_t i, size\_t j, bool check\_bounds=true)

#### Column/row wise interchange

- void swap rows (size t i0, size t i1, bool check bounds=true)
- void swap cols (size t j0, size t j1, bool check bounds=true)
- void zero\_row (size\_t i, bool check\_bounds=true)
- void zero\_col (size\_t j, bool check\_bounds=true)

#### **Arithmetic operators**

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

## **Public Attributes**

• bool visited = false

#### **Friends**

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

## 8.4.1 Detailed Description

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

Baseline class for binary arrays.

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

## **Template Parameters**

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

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

## 8.4.2 Constructor & Destructor Documentation

## 8.4.2.1 BArrayDense() [1/6]

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

Zero-size array.

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

## 8.4.2.2 BArrayDense() [2/6]

Empty array.

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

# 8.4.2.3 BArrayDense() [3/6]

Edgelist with data.

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

## 8.4.2.4 BArrayDense() [4/6]

Edgelist with no data (simpler)

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

## 8.4.2.5 BArrayDense() [5/6]

Copy constructor.

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

## 8.4.2.6 BArrayDense() [6/6]

Move operator.

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

## 8.4.2.7 ~BArrayDense()

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

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

## 8.4.3 Member Function Documentation

#### 8.4.3.1 clear()

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

Definition at line 909 of file barraydense-meat.hpp.

#### 8.4.3.2 col() [1/2]

Definition at line 502 of file barraydense-meat.hpp.

## 8.4.3.3 col() [2/2]

Definition at line 488 of file barraydense-meat.hpp.

# 8.4.3.4 colsum()

Definition at line 1012 of file barraydense-meat.hpp.

## 8.4.3.5 D() [1/2]

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

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

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

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

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

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

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

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

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

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

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

## 8.4.3.9 default\_val()

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

Definition at line 575 of file barraydense-meat.hpp.

#### 8.4.3.10 get\_cell()

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

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

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

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

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

## 8.4.3.13 get\_data()

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

Definition at line 1002 of file barraydense-meat.hpp.

#### 8.4.3.14 get\_entries()

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

Get the edgelist.

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

Returns

Entries<Cell\_Type>

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

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

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

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

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

#### 8.4.3.17 insert cell() [1/2]

Definition at line 701 of file barraydense-meat.hpp.

## 8.4.3.18 insert\_cell() [2/2]

Definition at line 667 of file barraydense-meat.hpp.

## 8.4.3.19 is\_dense()

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

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

## 8.4.3.20 is\_empty()

Definition at line 543 of file barraydense-meat.hpp.

## 8.4.3.21 ncol()

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

Definition at line 560 of file barraydense-meat.hpp.

# 8.4.3.22 nnozero()

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

Definition at line 564 of file barraydense-meat.hpp.

#### 8.4.3.23 nrow()

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

Definition at line 556 of file barraydense-meat.hpp.

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

Definition at line 619 of file barraydense-meat.hpp.

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

Definition at line 630 of file barraydense-meat.hpp.

## 8.4.3.26 operator\*=()

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

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

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

Definition at line 580 of file barraydense-meat.hpp.

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

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

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

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

# 8.4.3.33 operator/=()

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

Move assignment.

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

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

Assignment constructor.

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

## 8.4.3.36 operator==()

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

#### 8.4.3.37 out\_of\_range()

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

### 8.4.3.38 print()

Definition at line 968 of file barraydense-meat.hpp.

#### 8.4.3.39 reserve()

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

Definition at line 959 of file barraydense-meat.hpp.

#### 8.4.3.40 resize()

Definition at line 923 of file barraydense-meat.hpp.

## 8.4.3.41 rm\_cell()

Definition at line 644 of file barraydense-meat.hpp.

# 8.4.3.42 row() [1/2]

Definition at line 474 of file barraydense-meat.hpp.

## 8.4.3.43 row() [2/2]

Definition at line 461 of file barraydense-meat.hpp.

#### 8.4.3.44 rowsum()

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

## 8.4.3.45 set\_data()

Set the data object.

#### **Parameters**

data_	
delete_⊸	
data_	

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

## 8.4.3.46 swap\_cells()

```
template<typename Cell_Type , typename Data_Type >
void BArrayDense< Cell_Type, Data_Type >::swap_cells (
    size_t i0,
    size_t j0,
    size_t i1,
    size_t j1,
    bool check_bounds = true,
    int check_exists = CHECK::BOTH,
    int * report = nullptr ) [inline]
```

Definition at line 732 of file barraydense-meat.hpp.

### 8.4.3.47 swap\_cols()

Definition at line 816 of file barraydense-meat.hpp.

#### 8.4.3.48 swap\_rows()

Definition at line 788 of file barraydense-meat.hpp.

## 8.4.3.49 toggle\_cell()

Definition at line 769 of file barraydense-meat.hpp.

## 8.4.3.50 toggle\_lock()

#### 8.4.3.51 transpose()

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

Definition at line 881 of file barraydense-meat.hpp.

## 8.4.3.52 zero\_col()

Definition at line 862 of file barraydense-meat.hpp.

#### 8.4.3.53 zero\_row()

Definition at line 843 of file barraydense-meat.hpp.

#### 8.4.4 Friends And Related Function Documentation

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

#### 8.4.5 Member Data Documentation

#### 8.4.5.1 visited

```
template<typename Cell_Type = bool, typename Data_Type = bool>
bool BArrayDense< Cell_Type, Data_Type >::visited = false
```

This is as a reference, if we need to iterate through the cells and we need to keep track which were visited, we use this as a reference. So that if cell.visited = true and visited = true, it means that we haven't been here yet. Ideally, any routine using this->visited should switch it at the beginning of the routine.

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

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

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

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

#include <barraydensecell-bones.hpp>

#### **Public Member Functions**

- BArrayDenseCell (BArrayDense< Cell\_Type, Data\_Type > \*Array\_, size\_t i\_, size\_t j\_, bool check\_←
  bounds=true)
- BArrayDenseCell< Cell\_Type, Data\_Type > & operator= (const BArrayDenseCell< Cell\_Type, Data\_Type > &other)
- ∼BArrayDenseCell ()
- void operator= (const Cell Type &val)
- void operator+= (const Cell\_Type &val)
- void operator-= (const Cell\_Type &val)
- void operator\*= (const Cell\_Type &val)
- void operator/= (const Cell\_Type &val)
- operator Cell Type () const
- bool operator== (const Cell\_Type &val) const

#### **Friends**

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

## 8.5.1 Detailed Description

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

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

## 8.5.2 Constructor & Destructor Documentation

## 8.5.2.1 BArrayDenseCell()

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

## 8.5.2.2 ~BArrayDenseCell()

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

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

## 8.5.3 Member Function Documentation

## 8.5.3.1 operator Cell\_Type()

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

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

#### 8.5.3.2 operator\*=()

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

#### 8.5.3.3 operator+=()

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

#### 8.5.3.4 operator-=()

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

## 8.5.3.5 operator/=()

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

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

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

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

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

#### 8.5.3.8 operator==()

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

#### 8.5.4 Friends And Related Function Documentation

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

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

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

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

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

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

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

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

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

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

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

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

## 8.6.1 Detailed Description

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

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

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

• include/barry/barraydense-bones.hpp

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

#include <barraydensecol-bones.hpp>

#### **Public Member Functions**

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

## **Friends**

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

## 8.7.1 Detailed Description

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

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

### 8.7.2 Constructor & Destructor Documentation

#### 8.7.2.1 BArrayDenseCol()

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

#### 8.7.3 Member Function Documentation

## 8.7.3.1 begin()

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

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

#### 8.7.3.2 end()

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

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

## 8.7.3.3 operator()()

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

## 8.7.3.4 size()

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

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

## 8.7.4 Friends And Related Function Documentation

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

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

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

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

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

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

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

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

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

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

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

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

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

### **Public Member Functions**

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

#### **Friends**

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

## 8.8.1 Detailed Description

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

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

## 8.8.2 Constructor & Destructor Documentation

#### 8.8.2.1 BArrayDenseCol\_const()

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

# 8.8.3 Member Function Documentation

## 8.8.3.1 begin()

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

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

## 8.8.3.2 end()

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

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

#### 8.8.3.3 operator()()

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

#### 8.8.3.4 size()

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

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

#### 8.8.4 Friends And Related Function Documentation

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

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

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

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

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

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

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

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

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

#include <barraydenserow-bones.hpp>

# **Public Member Functions**

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

#### Friends

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

## 8.9.1 Detailed Description

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

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

#### 8.9.2 Constructor & Destructor Documentation

### 8.9.2.1 BArrayDenseRow()

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

## 8.9.3 Member Function Documentation

## 8.9.3.1 begin()

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

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

## 8.9.3.2 end()

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

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

#### 8.9.3.3 operator()()

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

#### 8.9.3.4 size()

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

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

## 8.9.4 Friends And Related Function Documentation

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

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

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

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

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

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

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

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

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

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

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

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

#include <barraydenserow-bones.hpp>

#### **Public Member Functions**

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

#### **Friends**

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

## 8.10.1 Detailed Description

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

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

## 8.10.2 Constructor & Destructor Documentation

#### 8.10.2.1 BArrayDenseRow\_const()

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

## 8.10.3 Member Function Documentation

#### 8.10.3.1 begin()

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

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

## 8.10.3.2 end()

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

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

#### 8.10.3.3 operator()()

```
template<typename Cell_Type = bool, typename Data_Type = bool> const std::pair<size_t,Cell<Cell_Type> > BArrayDenseRow_const< Cell_Type, Data_Type > \leftrightarrow ::operator() ( size_t i ) const [inline]
```

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

### 8.10.3.4 size()

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

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

## 8.10.4 Friends And Related Function Documentation

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

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

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

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

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

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

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

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

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

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

#### **Public Member Functions**

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

## 8.11.1 Detailed Description

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

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

## 8.11.2 Constructor & Destructor Documentation

## 8.11.2.1 BArrayRow()

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

#### 8.11.2.2 $\sim$ BArrayRow()

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

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

## 8.11.3 Member Function Documentation

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

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

## 8.11.3.2 operator\*=()

# 8.11.3.3 operator+=()

#### 8.11.3.4 operator-=()

#### 8.11.3.5 operator/=()

#### 8.11.3.6 operator=()

## 8.11.3.7 operator==()

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

• include/barry/barrayrow-bones.hpp

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

#include <barrayrow-bones.hpp>

# **Public Member Functions**

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

## 8.12.1 Detailed Description

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

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

## 8.12.2 Constructor & Destructor Documentation

#### 8.12.2.1 BArrayRow\_const()

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

#### 8.12.2.2 ~BArrayRow\_const()

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

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

#### 8.12.3 Member Function Documentation

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

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

## 8.12.3.2 operator"!=()

#### 8.12.3.3 operator<()

#### 8.12.3.4 operator<=()

#### 8.12.3.5 operator==()

# 8.12.3.6 operator>()

## 8.12.3.7 operator>=()

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

• include/barry/barrayrow-bones.hpp

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

Row or column of a BArray

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

## **Public Member Functions**

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

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

## 8.13.1 Detailed Description

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

Row or column of a BArray

**Template Parameters** 

Cell_Type	
Data_Type	

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

## 8.13.2 Constructor & Destructor Documentation

## 8.13.2.1 BArrayVector()

Construct a new BArrayVector object.

## **Parameters**

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

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

#### 8.13.2.2 ~BArrayVector()

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

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

#### 8.13.3 Member Function Documentation

## 8.13.3.1 begin()

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

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

#### 8.13.3.2 end()

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

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

# 8.13.3.3 is\_col()

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

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

## 8.13.3.4 is\_row()

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

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

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

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

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

## 8.13.3.6 operator \*= ()

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

#### 8.13.3.7 operator+=()

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

#### 8.13.3.8 operator-=()

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

#### 8.13.3.9 operator/=()

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

## 8.13.3.10 operator=()

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

## 8.13.3.11 operator==()

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

#### 8.13.3.12 size()

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

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

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

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

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

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

#### **Public Member Functions**

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

## 8.14.1 Detailed Description

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

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

#### 8.14.2 Constructor & Destructor Documentation

# 8.14.2.1 BArrayVector\_const()

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

#### 8.14.2.2 ~BArrayVector\_const()

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

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

## 8.14.3 Member Function Documentation

## 8.14.3.1 begin()

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

## 8.14.3.2 end()

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

## 8.14.3.3 is\_col()

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

## 8.14.3.4 is\_row()

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

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

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

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

#### 8.14.3.6 operator"!=()

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

## 8.14.3.7 operator<()

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

## 8.14.3.8 operator<=()

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

# 8.14.3.9 operator==()

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

## 8.14.3.10 operator>()

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

#### 8.14.3.11 operator>=()

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

## 8.14.3.12 size()

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

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

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

# 8.15 Cell< Cell\_Type > Class Template Reference

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

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

#### **Public Member Functions**

- Cell ()
- Cell (Cell\_Type value\_, bool visited\_=false, bool active\_=true)
- ~Cell ()
- Cell (const Cell< Cell\_Type > &arg)
- Cell< Cell\_Type > & operator= (const Cell< Cell\_Type > &other)
- Cell (Cell< Cell\_Type > &&arg) noexcept
- Cell< Cell\_Type > & operator= (Cell< Cell\_Type > &&other) noexcept
- void add (Cell\_Type x)
- operator Cell\_Type () const
- bool operator== (const Cell< Cell\_Type > &rhs) const
- bool operator!= (const Cell< Cell\_Type > &rhs) const
- void add (double x)
- void add (size\_t x)
- void add (int x)
- Cell ()
- Cell ()
- Cell ()

# **Public Attributes**

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

# 8.15.1 Detailed Description

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

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

· value: the content

· visited: boolean (just a convenient)

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

#### 8.15.2 Constructor & Destructor Documentation

## 8.15.2.1 Cell() [1/7]

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

# 8.15.2.2 Cell() [2/7]

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

## 8.15.2.3 $\sim$ Cell()

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

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

## 8.15.2.4 Cell() [3/7]

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

## 8.15.2.5 Cell() [4/7]

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

## 8.15.2.6 Cell() [5/7]

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

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

## 8.15.2.7 Cell() [6/7]

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

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

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

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

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

## 8.15.3 Member Function Documentation

## 8.15.3.1 add() [1/4]

## 8.15.3.2 add() [2/4]

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

## 8.15.3.3 add() [3/4]

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

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

## 8.15.3.4 add() [4/4]

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

## 8.15.3.5 operator Cell\_Type()

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

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

# 8.15.3.6 operator"!=()

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

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

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

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

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

## 8.15.3.9 operator==()

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

## 8.15.4 Member Data Documentation

## 8.15.4.1 active

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

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

## 8.15.4.2 value

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

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

#### 8.15.4.3 visited

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

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

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

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

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

## 8.16.1 Detailed Description

```
template<typename Cell_Type> class Cell_const< Cell_Type>
```

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

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

• include/barry/barray-meat.hpp

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

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

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



## **Public Member Functions**

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

## **Public Attributes**

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

# 8.17.1 Detailed Description

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

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

## 8.17.2 Constructor & Destructor Documentation

# 8.17.2.1 ConstBArrayRowlter()

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

#### 8.17.2.2 ∼ConstBArrayRowlter()

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

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

#### 8.17.3 Member Data Documentation

#### 8.17.3.1 Array

```
template<typename Cell_Type , typename Data_Type >
const BArray<Cell_Type,Data_Type>* ConstBArrayRowIter< Cell_Type, Data_Type >::Array
```

Definition at line 15 of file barray-iterator.hpp.

#### 8.17.3.2 current\_col

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

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

#### 8.17.3.3 current\_row

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

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

#### 8.17.3.4 iter

```
template<typename Cell_Type , typename Data_Type >
Row_type<Cell_Type>::const_iterator ConstBArrayRowIter< Cell_Type, Data_Type >::iter
```

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

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

• include/barry/barray-iterator.hpp

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

A counter function based on change statistics.

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

## **Public Member Functions**

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

Creator passing a counter and an initializer

#### **Parameters**

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

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

Move constructor

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

Move assignment.

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

Get and set the hasher function.

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

## **Public Attributes**

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

# 8.18.1 Detailed Description

template<typename Array\_Type = BArray<>, typename Data\_Type = bool> class Counter< Array\_Type, Data\_Type >

A counter function based on change statistics.

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

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

## 8.18.2 Constructor & Destructor Documentation

## 8.18.2.1 Counter() [1/4]

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

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

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

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

# 8.18.2.3 Counter() [3/4]

Copy constructor.

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

Move constructor.

#### 8.18.2.5 ∼Counter()

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

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

#### 8.18.3 Member Function Documentation

## 8.18.3.1 count()

## 8.18.3.2 get\_description()

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

#### 8.18.3.3 get\_hasher()

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

#### 8.18.3.4 get\_name()

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

## 8.18.3.5 init()

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

Copy assignment.

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

Move assignment.

## 8.18.3.8 set\_hasher()

Get and set the hasher function.

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

**Parameters** 

fun

## 8.18.4 Member Data Documentation

## 8.18.4.1 count\_fun

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

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

#### 8.18.4.2 data

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

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

#### 8.18.4.3 desc

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

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

#### 8.18.4.4 hasher\_fun

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

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

## 8.18.4.5 init\_fun

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

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

## 8.18.4.6 name

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

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

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

• include/barry/counters-bones.hpp

# 8.19 Counters < Array\_Type, Data\_Type > Class Template Reference

Vector of counters.

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

#### **Public Member Functions**

- Counters ()
- ∼Counters ()
- Counters (const Counters < Array\_Type, Data\_Type > &counter\_)

Copy constructor.

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

Move constructor.

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

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

Move assignment constructor.

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

Returns a pointer to a particular counter.

• std::size\_t size () const noexcept

Number of counters in the set.

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

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

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

## 8.19.1 Detailed Description

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

Vector of counters.

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

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

#### 8.19.2 Constructor & Destructor Documentation

## 8.19.2.1 Counters() [1/3]

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

## 8.19.2.2 ∼Counters()

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

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

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

Copy constructor.

## **Parameters**



## 8.19.2.4 Counters() [3/3]

Move constructor.

#### **Parameters**

counters⊷

## 8.19.3 Member Function Documentation

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

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

#### 8.19.3.3 add hash()

## 8.19.3.4 gen\_hash()

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

#### **Parameters**

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

## Returns

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

#### 8.19.3.5 get\_descriptions()

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

# 8.19.3.6 get\_names()

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

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

Copy assignment constructor.

#### **Parameters**



#### Returns

Counters<Array\_Type,Data\_Type>

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

Move assignment constructor.

## **Parameters**



#### Returns

```
Counters<Array_Type,Data_Type>&
```

## 8.19.3.9 operator[]()

Returns a pointer to a particular counter.

#### **Parameters**

```
idx Id of the counter
```

#### Returns

```
Counter<Array_Type,Data_Type>*
```

## 8.19.3.10 size()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
std::size_t Counters< Array_Type, Data_Type >::size ( ) const [inline], [noexcept]
```

Number of counters in the set.

## Returns

size\_t

Definition at line 164 of file counters-bones.hpp.

The documentation for this class was generated from the following file:

• include/barry/counters-bones.hpp

8.20 DEFM Class Reference 139

# 8.20 DEFM Class Reference

#include <defm-bones.hpp>

Inheritance diagram for DEFM:



Collaboration diagram for DEFM:



## **Public Member Functions**

- DEFM (int \*id, int \*y, double \*x, size\_t id\_length, size\_t y\_ncol, size\_t x\_ncol, size\_t m\_order, bool copy\_
   data=true, bool column\_major=true)
- DEFMModel & get\_model ()
- void init ()
- void simulate (std::vector< double > par, int \*y\_out)
- size\_t get\_n\_y () const
- size\_t get\_n\_obs () const
- size\_t get\_n\_covars () const
- size\_t get\_m\_order () const
- size\_t get\_n\_rows () const
- const int \* get\_Y () const
- const int \* get\_ID () const
- const double \* get\_X () const
- barry::FreqTable< int > motif\_census (std::vector< size\_t > idx)

```
std::vector< double > logodds (const std::vector< double > &par, size_t i, size_t j)
void set_names (std::vector< std::string > Y_names_, std::vector< std::string > X_names_)
const std::vector< std::string > & get_Y_names () const
const std::vector< std::string > & get_X_names () const
void print () const
std::vector< bool > is_motif ()
bool get_column_major () const noexcept
```

## 8.20.1 Detailed Description

Definition at line 4 of file defm-bones.hpp.

## 8.20.2 Constructor & Destructor Documentation

#### 8.20.2.1 DEFM()

```
DEFM::DEFM (
        int * id,
        int * y,
        double * x,
        size_t id_length,
        size_t y_ncol,
        size_t x_ncol,
        size_t m_order,
        bool copy_data = true,
        bool column_major = true ) [inline]
```

Definition at line 105 of file defm-meat.hpp.

## 8.20.3 Member Function Documentation

```
8.20.3.1 get_column_major()
```

```
bool DEFM::get_column_major ( ) const [inline], [noexcept]
```

Definition at line 449 of file defm-meat.hpp.

# 8.20.3.2 get\_ID()

```
const int * DEFM::get_ID ( ) const [inline]
```

Definition at line 310 of file defm-meat.hpp.

8.20 DEFM Class Reference 141

## 8.20.3.3 get\_m\_order()

```
size_t DEFM::get_m_order ( ) const [inline]
```

Definition at line 295 of file defm-meat.hpp.

## 8.20.3.4 get\_model()

```
DEFMModel& DEFM::get_model ( ) [inline]
```

Definition at line 49 of file defm-bones.hpp.

## 8.20.3.5 get\_n\_covars()

```
size_t DEFM::get_n_covars ( ) const [inline]
```

Definition at line 290 of file defm-meat.hpp.

## 8.20.3.6 get\_n\_obs()

```
size_t DEFM::get_n_obs ( ) const [inline]
```

Definition at line 285 of file defm-meat.hpp.

## 8.20.3.7 get\_n\_rows()

```
size_t DEFM::get_n_rows ( ) const [inline]
```

Definition at line 300 of file defm-meat.hpp.

# 8.20.3.8 get\_n\_y()

```
size_t DEFM::get_n_y ( ) const [inline]
```

Definition at line 280 of file defm-meat.hpp.

## 8.20.3.9 get\_X()

```
const double * DEFM::get_X ( ) const [inline]
```

Definition at line 315 of file defm-meat.hpp.

## 8.20.3.10 get\_X\_names()

```
\verb|const| std::vector<| std::string| > \& DEFM::get_X_names ( ) | const| [inline]|
```

Definition at line 422 of file defm-meat.hpp.

## 8.20.3.11 get\_Y()

```
const int * DEFM::get_Y ( ) const [inline]
```

Definition at line 305 of file defm-meat.hpp.

# 8.20.3.12 get\_Y\_names()

```
const std::vector< std::string > & DEFM::get_Y_names ( ) const [inline]
```

Definition at line 418 of file defm-meat.hpp.

## 8.20.3.13 init()

```
void DEFM::init ( ) [inline]
```

Definition at line 215 of file defm-meat.hpp.

# 8.20.3.14 is\_motif()

```
std::vector< bool > DEFM::is_motif ( ) [inline]
```

Definition at line 439 of file defm-meat.hpp.

8.20 DEFM Class Reference 143

#### 8.20.3.15 logodds()

Definition at line 359 of file defm-meat.hpp.

## 8.20.3.16 motif\_census()

```
barry::FreqTable< int > DEFM::motif_census (  std::vector < size_t > \mathit{idx} ) \quad [inline]
```

Definition at line 321 of file defm-meat.hpp.

#### 8.20.3.17 print()

```
void DEFM::print ( ) const [inline]
```

Definition at line 426 of file defm-meat.hpp.

## 8.20.3.18 set\_names()

```
void DEFM::set_names ( std::vector < std::string > \textit{Y}\_names\_, \\ std::vector < std::string > \textit{X}\_names\_ ) \quad [inline]
```

Definition at line 401 of file defm-meat.hpp.

## 8.20.3.19 simulate()

```
void DEFM::simulate (
          std::vector< double > par,
          int * y_out ) [inline]
```

Definition at line 39 of file defm-meat.hpp.

The documentation for this class was generated from the following files:

- include/barry/models/defm/defm-bones.hpp
- include/barry/models/defm/defm-meat.hpp

## 8.21 DEFMCounterData Class Reference

Data class used to store arbitrary size\_t or double vectors.

```
#include <defm-types.hpp>
```

#### **Public Member Functions**

- DEFMCounterData ()
- DEFMCounterData (const std::vector< size\_t > indices\_, const std::vector< double > numbers\_, const std::vector< bool > logical\_, bool is\_motif\_=true)
- size\_t idx (size\_t i) const
- double num (size\_t i) const
- · bool is true (size ti) const
- ∼DEFMCounterData ()

## **Public Attributes**

- std::vector< size\_t > indices
- std::vector< double > numbers
- std::vector< bool > logical
- · bool is\_motif

If false, then is a logit intercept.

## 8.21.1 Detailed Description

Data class used to store arbitrary size\_t or double vectors.

Definition at line 64 of file defm-types.hpp.

The documentation for this class was generated from the following file:

• include/barry/models/defm/defm-types.hpp

## 8.22 DEFMData Class Reference

Data class for **DEFM** arrays.

```
#include <defm-types.hpp>
```

#### **Public Member Functions**

• DEFMData ()

Vector indicating which covariates are included in the model.

DEFMData (DEFMArray \*array\_, const double \*covariates\_, size\_t obs\_start\_, size\_t X\_ncol\_, size\_t X\_← nrow )

Constructor.

double operator() (size t i, size t j) const

Access to the row (i) colum (j) data.

- double at (size\_t i, size\_t j) const
- size\_t ncol () const
- size\_t nrow () const
- · void print () const
- ∼DEFMData ()

#### **Public Attributes**

- DEFMArray \* array
- const double \* covariates

Vector of covariates (complete vector)

size\_t obs\_start

Index of the observation in the data.

size\_t X\_ncol

Number of columns in the array of covariates.

size\_t X\_nrow

Number of rows in the array of covariates.

- std::vector< size\_t > covar\_sort
- std::vector< size\_t > covar\_used

Value where the sorting of the covariates is stored.

#### 8.22.1 Detailed Description

Data class for **DEFM** arrays.

This holds information pointing to the data array, including information regarding the number of observations, the time slices of the observation, and the number of covariates in the data.

Definition at line 16 of file defm-types.hpp.

The documentation for this class was generated from the following file:

• include/barry/models/defm/defm-types.hpp

# 8.23 DEFMRuleData Class Reference

#include <defm-types.hpp>

Inheritance diagram for DEFMRuleData:



#### **Public Member Functions**

- double num (size\_t i) const
- size\_t idx (size\_t i) const
- bool is\_true (size\_t i) const
- DEFMRuleData ()
- DEFMRuleData (std::vector< double > numbers\_, std::vector< size\_t > indices\_, std::vector< bool > logical\_)
- DEFMRuleData (std::vector< double > numbers\_, std::vector< size\_t > indices\_)

## **Public Attributes**

- std::vector< double > numbers
- std::vector< size\_t > indices
- std::vector< bool > logical
- bool init = false

## 8.23.1 Detailed Description

Definition at line 89 of file defm-types.hpp.

The documentation for this class was generated from the following file:

• include/barry/models/defm/defm-types.hpp

# 8.24 DEFMRuleDynData Class Reference

#include <defm-types.hpp>

Inheritance diagram for DEFMRuleDynData:



Collaboration diagram for DEFMRuleDynData:



## **Public Member Functions**

- DEFMRuleDynData (const std::vector< double > \*counts\_, std::vector< double > numbers\_={}, std::vector< size\_t > indices\_={}, std::vector< bool > logical\_={})
- ∼DEFMRuleDynData ()

## **Public Attributes**

const std::vector< double > \* counts

# 8.24.1 Detailed Description

Definition at line 152 of file defm-types.hpp.

The documentation for this class was generated from the following file:

include/barry/models/defm/defm-types.hpp

# 8.25 Entries < Cell\_Type > Class Template Reference

A wrapper class to store source, target, val from a BArray object.

```
#include <typedefs.hpp>
```

#### **Public Member Functions**

- Entries ()
- Entries (size\_t n)
- ∼Entries ()
- void resize (size\_t n)

## **Public Attributes**

```
• std::vector< size_t > source
```

- std::vector< size\_t > target
- std::vector< Cell\_Type > val

#### 8.25.1 Detailed Description

```
\label{lem:continuous} \begin{split} \text{template} &< \text{typename Cell\_Type} > \\ \text{class Entries} &< \text{Cell\_Type} > \end{split}
```

A wrapper class to store source, target, val from a BArray object.

**Template Parameters** 

```
Cell_Type Any type
```

Definition at line 78 of file typedefs.hpp.

#### 8.25.2 Constructor & Destructor Documentation

#### 8.25.2.1 Entries() [1/2]

```
template<typename Cell_Type >
Entries< Cell_Type >::Entries ( ) [inline]
```

Definition at line 84 of file typedefs.hpp.

#### 8.25.2.2 Entries() [2/2]

Definition at line 85 of file typedefs.hpp.

# 8.25.2.3 $\sim$ Entries()

```
template<typename Cell_Type >
Entries< Cell_Type >::~Entries ( ) [inline]
```

Definition at line 92 of file typedefs.hpp.

#### 8.25.3 Member Function Documentation

#### 8.25.3.1 resize()

Definition at line 94 of file typedefs.hpp.

#### 8.25.4 Member Data Documentation

#### 8.25.4.1 source

```
template<typename Cell_Type >
std::vector< size_t > Entries< Cell_Type >::source
```

Definition at line 80 of file typedefs.hpp.

# 8.25.4.2 target

```
template<typename Cell_Type >
std::vector< size_t > Entries< Cell_Type >::target
```

Definition at line 81 of file typedefs.hpp.

#### 8.25.4.3 val

```
template<typename Cell_Type >
std::vector< Cell_Type > Entries< Cell_Type >::val
```

Definition at line 82 of file typedefs.hpp.

The documentation for this class was generated from the following file:

• include/barry/typedefs.hpp

## 8.26 Flock Class Reference

A Flock is a group of Geese.

```
#include <flock-bones.hpp>
```

#### **Public Member Functions**

- Flock ()
- ∼Flock ()
- size\_t add\_data (std::vector< std::vector< size\_t > &annotations, std::vector< size\_t > &geneid, std
   ::vector< int > &parent, std::vector< bool > &duplication)

Add a tree to the flock.

• void set seed (const size t &s)

Set the seed of the model.

- void init (size\_t bar\_width=BARRY\_PROGRESS\_BAR\_WIDTH)
- PhyloCounters \* get\_counters ()
- PhyloSupport \* get support fun ()
- std::vector< double > \* get stats support ()
- std::vector< std::vector< double > > \* get\_stats\_target ()
- PhyloModel \* get\_model ()

Returns the joint likelihood of the model.

Geese \* operator() (size\_t i, bool check\_bounds=true)

Access the i-th geese element.

#### Information about the model

- size\_t nfuns () const noexcept
- size\_t ntrees () const noexcept
- std::vector< size\_t > nnodes () const noexcept
- std::vector< size\_t > nleafs () const noexcept
- size\_t nterms () const
- size\_t support\_size () const noexcept
- std::vector< std::string > colnames () const
- $\bullet \ \ \mathsf{size\_t} \ \mathsf{parse\_polytomies} \ (\mathsf{bool} \ \mathsf{verb=true}, \ \mathsf{std} :: \mathsf{vector} < \ \mathsf{size\_t} > * \mathsf{dist=nullptr}) \ \mathsf{const} \ \mathsf{noexcept}$

Check polytomies and return the largest.

void print () const

8.26 Flock Class Reference 151

## **Public Attributes**

- std::vector< Geese > dat
- size\_t nfunctions = 0u
- bool initialized = false
- std::mt19937 rengine
- PhyloModel model = PhyloModel()

## 8.26.1 Detailed Description

A Flock is a group of Geese.

This object buils a model with multiple trees (Geese objects), with all of these using the same PhyloModel object. Available counters (terms) can be found in counter-phylo.

Definition at line 14 of file flock-bones.hpp.

#### 8.26.2 Constructor & Destructor Documentation

## 8.26.2.1 Flock()

```
Flock::Flock ( ) [inline]
```

Definition at line 25 of file flock-bones.hpp.

## 8.26.2.2 ∼Flock()

```
Flock::~Flock ( ) [inline]
```

Definition at line 26 of file flock-bones.hpp.

## 8.26.3 Member Function Documentation

## 8.26.3.1 add\_data()

Add a tree to the flock.

#### **Parameters**

annotations	see Geese::Geese.
geneid	see Geese.
parent	see Geese.
duplication	see Geese.

#### Returns

size\_t The number of tree in the model (starting from zero).

Definition at line 6 of file flock-meat.hpp.

#### 8.26.3.2 colnames()

```
std::vector< std::string > Flock::colnames ( ) const [inline]
```

Definition at line 246 of file flock-meat.hpp.

## 8.26.3.3 get\_counters()

```
PhyloCounters * Flock::get_counters ( ) [inline]
```

Definition at line 100 of file flock-meat.hpp.

## 8.26.3.4 get\_model()

```
PhyloModel * Flock::get_model ( ) [inline]
```

Definition at line 131 of file flock-meat.hpp.

# 8.26.3.5 get\_stats\_support()

```
\verb|std::vector<| double > * Flock::get_stats_support () | [inline]|
```

Definition at line 117 of file flock-meat.hpp.

8.26 Flock Class Reference 153

## 8.26.3.6 get\_stats\_target()

```
\verb|std::vector| < \verb|std::vector| < \verb|double| > > * Flock::get_stats_target () | [inline]|
```

Definition at line 124 of file flock-meat.hpp.

#### 8.26.3.7 get\_support\_fun()

```
PhyloSupport * Flock::get_support_fun ( ) [inline]
```

Definition at line 110 of file flock-meat.hpp.

## 8.26.3.8 init()

Definition at line 49 of file flock-meat.hpp.

#### 8.26.3.9 likelihood\_joint()

Returns the joint likelihood of the model.

## Parameters

par	Vector of model parameters.	
as_log	When true it will return the value as log.	
use_reduced_sequence	When true (default) will compute the likelihood using the reduced sequence,	
	which is faster.	

#### Returns

double

Definition at line 138 of file flock-meat.hpp.

#### 8.26.3.10 nfuns()

```
size_t Flock::nfuns ( ) const [inline], [noexcept]
```

Definition at line 189 of file flock-meat.hpp.

## 8.26.3.11 nleafs()

```
std::vector< size_t > Flock::nleafs ( ) const [inline], [noexcept]
```

Definition at line 217 of file flock-meat.hpp.

#### 8.26.3.12 nnodes()

```
std::vector< size_t > Flock::nnodes ( ) const [inline], [noexcept]
```

Definition at line 203 of file flock-meat.hpp.

# 8.26.3.13 nterms()

```
size_t Flock::nterms ( ) const [inline]
```

Definition at line 231 of file flock-meat.hpp.

## 8.26.3.14 ntrees()

```
size_t Flock::ntrees ( ) const [inline], [noexcept]
```

Definition at line 196 of file flock-meat.hpp.

## 8.26.3.15 operator()()

Access the i-th geese element.

8.26 Flock Class Reference 155

#### **Parameters**

i	Element to access
check_bounds	When true, it will check bounds.

Returns

Geese \*

Definition at line 324 of file flock-meat.hpp.

# 8.26.3.16 parse\_polytomies()

Check polytomies and return the largest.

Definition at line 253 of file flock-meat.hpp.

## 8.26.3.17 print()

```
void Flock::print ( ) const [inline]
```

Definition at line 280 of file flock-meat.hpp.

# 8.26.3.18 set\_seed()

Set the seed of the model.

**Parameters** 

```
s Passed to the rengine.seed() member object.
```

Definition at line 42 of file flock-meat.hpp.

## 8.26.3.19 support\_size()

```
size_t Flock::support_size ( ) const [inline], [noexcept]
```

Definition at line 239 of file flock-meat.hpp.

## 8.26.4 Member Data Documentation

## 8.26.4.1 dat

```
std::vector< Geese > Flock::dat
```

Definition at line 17 of file flock-bones.hpp.

## 8.26.4.2 initialized

```
bool Flock::initialized = false
```

Definition at line 19 of file flock-bones.hpp.

#### 8.26.4.3 model

```
PhyloModel Flock::model = PhyloModel()
```

Definition at line 23 of file flock-bones.hpp.

## 8.26.4.4 nfunctions

```
size_t Flock::nfunctions = 0u
```

Definition at line 18 of file flock-bones.hpp.

#### 8.26.4.5 rengine

```
std::mt19937 Flock::rengine
```

Definition at line 22 of file flock-bones.hpp.

The documentation for this class was generated from the following files:

- include/barry/models/geese/flock-bones.hpp
- include/barry/models/geese/flock-meat.hpp

# 8.27 FreqTable < T > Class Template Reference

Frequency table of vectors.

```
#include <freqtable.hpp>
```

#### **Public Member Functions**

```
• FreqTable ()
```

- ∼FreqTable ()
- size\_t add (const std::vector< T > &x, size\_t \*h\_precomp)
- Counts\_type as\_vector () const
- const std::vector< double > & get\_data () const
- const std::unordered\_map< size\_t, size\_t > & get\_index () const
- void clear ()
- void reserve (size\_t n, size\_t k)
- void print () const
- size\_t size () const noexcept

Number of unique elements in the table. (.

size\_t make\_hash (const std::vector< T > &x) const

# 8.27.1 Detailed Description

```
template<typename T = double> class FreqTable< T >
```

Frequency table of vectors.

This is mostly used in Support. The main data is contained in the data double vector. The matrix is stored in a row-wise fashion, where the first element is the frequency with which the vector is observed.

For example, in a model with k terms the first k+1 elements of data would be:

- · weights
- term 1
- term 2
- ...
- · term k

Definition at line 22 of file freqtable.hpp.

## 8.27.2 Constructor & Destructor Documentation

## 8.27.2.1 FreqTable()

```
template<typename T = double>
FreqTable< T >::FreqTable ( ) [inline]
```

Definition at line 34 of file freqtable.hpp.

#### 8.27.2.2 ∼FreqTable()

```
template<typename T = double>
FreqTable< T >::~FreqTable ( ) [inline]
```

Definition at line 35 of file freqtable.hpp.

#### 8.27.3 Member Function Documentation

# 8.27.3.1 add()

Definition at line 59 of file freqtable.hpp.

## 8.27.3.2 as\_vector()

```
template<typename T >
Counts_type FreqTable< T >::as_vector [inline]
```

Definition at line 139 of file freqtable.hpp.

#### 8.27.3.3 clear()

```
template<typename T >
void FreqTable< T >::clear [inline]
```

Definition at line 168 of file freqtable.hpp.

#### 8.27.3.4 get\_data()

```
template<typename T = double>
const std::vector< double >& FreqTable< T >::get_data ( ) const [inline]
```

Definition at line 40 of file freqtable.hpp.

## 8.27.3.5 get\_index()

```
template<typename T = double>
const std::unordered_map<size_t,size_t>& FreqTable< T >::get_index ( ) const [inline]
```

Definition at line 41 of file freqtable.hpp.

#### 8.27.3.6 make\_hash()

Definition at line 239 of file freqtable.hpp.

# 8.27.3.7 print()

```
template<typename T >
void FreqTable< T >::print [inline]
```

Definition at line 204 of file freqtable.hpp.

#### 8.27.3.8 reserve()

Definition at line 182 of file freqtable.hpp.

#### 8.27.3.9 size()

```
template<typename T >
size_t FreqTable< T >::size [inline], [noexcept]
```

Number of unique elements in the table. (.

Returns

size t

Definition at line 231 of file freqtable.hpp.

The documentation for this class was generated from the following file:

• include/barry/freqtable.hpp

#### 8.28 Geese Class Reference

Annotated Phylo Model.

```
#include <geese-bones.hpp>
```

## **Public Member Functions**

- ∼Geese ()
- void init (size\_t bar\_width=BARRY\_PROGRESS\_BAR\_WIDTH)
- void inherit\_support (const Geese &model\_, bool delete\_support\_=false)
- void calc\_sequence (Node \*n=nullptr)
- void calc reduced sequence ()
- double likelihood (const std::vector< double > &par, bool as\_log=false, bool use\_reduced\_sequence=true, size\_t ncores=1u, bool no\_update\_pset\_probs=false)
- double likelihood\_exhaust (const std::vector< double > &par)
- std::vector< double > get\_probabilities () const
- void set\_seed (const size\_t &s)
- std::vector< std::vector< size\_t >> simulate (const std::vector< double > &par)
- std::vector< std::vector< double >> observed\_counts ()
- void print\_observed\_counts ()
- void print () const

8.28 Geese Class Reference 161

Prints information about the GEESE.

- void print\_nodes () const
- void init node (Node &n)
- void update annotations (size t nodeid, std::vector< size t > newann)
- std::vector< std::vector< bool >> get\_states () const

Powerset of a gene's possible states.

std::vector< size\_t > get\_annotated\_nodes () const

Returns the ids of the nodes with at least one annotation.

std::vector< size t > get annotations () const

Returns the annotations of the nodes with at least one annotation.

#### Construct a new Geese object

The model includes a total of N + 1 nodes, the + 1 beign the root node.

#### **Parameters**

annotations	A vector of vectors with annotations. It should be of length $k$ (number of functions). Each vector should be of length $\mathbb N$ (equal to the number of nodes, including interior). Possible values are 0, 1, and 9.
geneid	Id of the gene. It should be of length $\mathbb{N}$ .
parent	Id of the parent gene. Also of length N
duplication	Logical scalar indicating the type of event (true: duplication, false: speciation.)

The ordering of the entries does not matter. Passing the nodes in post order or not makes no difference to the constructor.

- Geese ()
- Geese (std::vector< std::vector< size\_t > &annotations, std::vector< size\_t > &geneid, std::vector< int > &parent, std::vector< bool > &duplication)
- Geese (const Geese &model\_, bool copy\_data=true)
- Geese (Geese &&x) noexcept
- Geese & operator= (const Geese &model\_)=delete
- Geese & operator= (Geese &&model\_) noexcept=delete

#### Information about the model

#### **Parameters**

verb When true it will print out information about the encountered polytomies.

size\_t nfuns () const noexcept

Number of functions analyzed.

size\_t nnodes () const noexcept

Number of nodes (interior + leaf)

• size\_t nleafs () const noexcept

Number of leaf.

• size t nterms () const

Number of terms included.

• size\_t support\_size () const noexcept

Number of unique sets of sufficient stats.

std::vector< size\_t > nannotations () const noexcept

Number of annotations.

std::vector< std::string > colnames () const

Names of the terms in the model.

• size\_t parse\_polytomies (bool verb=true, std::vector< size\_t > \*dist=nullptr) const noexcept

Check polytomies and return the largest.

#### Geese prediction

Calculate the conditional probability

#### **Parameters**

par	Vector of parameters (terms + root).	
res_prob	Vector indicating each nodes' state probability.	
leave_one_out	When true, it will compute the predictions using leave-one-out, thus the prediction will be repeated nleaf times.	
only_annotated	When true, it will make the predictions only on the induced sub-tree with annotated leafs.	
use_reduced_sequence	Passed to the likelihood method.	
preorder	For the tree traversal.	

When res\_prob is specified, the function will attach the member vector probabilities from the Nodes objects. This contains the probability that the ith node has either of the possible states.

#### Returns

std::vector< double > Returns the posterior probability

- std::vector< std::vector< double >> predict (const std::vector< double > &par, std::vector< std::vector< double >> \*res\_prob=nullptr, bool leave\_one\_out=false, bool only\_annotated=false, bool use\_reduced
  \_sequence=true)
- std::vector< std::vector< double > > predict\_backend (const std::vector< double > &par, bool use\_←
  reduced sequence, const std::vector< size t > &preorder)
- std::vector< std::vector< double > > predict\_exhaust\_backend (const std::vector< double > &par, const std::vector< size\_t > &preorder)
- std::vector< std::vector< double > > predict\_exhaust (const std::vector< double > &par)
- std::vector< std::vector< double > > predict\_sim (const std::vector< double > &par, bool only\_

   annotated=false, size\_t nsims=10000u)

## Non-const pointers to shared objects in <tt>Geese</tt>

These functions provide direct access to some member objects that are shared by the nodes within Geese.

#### Returns

```
get_rengine() returns the Pseudo-RNG engine used.
get_counters() returns the vector of counters used.
get_model() returns the Model object used.
get_support_fun() returns the computed support of the model.
```

- std::mt19937 \* get rengine ()
- PhyloCounters \* get\_counters ()
- PhyloModel \* get\_model ()
- PhyloSupport \* get support fun ()

#### **Public Attributes**

- size t nfunctions
- std::map< size t, Node > nodes
- barry::MapVec\_type< size\_t > map\_to\_state\_id

8.28 Geese Class Reference 163

```
    std::vector< std::vector< size_t >> > pset_loc
    Locations of columns.
```

- std::vector< size t > sequence
- std::vector< size\_t > reduced\_sequence
- bool initialized = false
- bool delete\_rengine = false
- bool delete\_support = false

#### **Static Public Attributes**

```
    static const size_t etype_default = 1ul
```

- static const size\_t etype\_speciation = 0ul
- static const size t etype duplication = 1ul
- static const size\_t etype\_either = 2ul

# 8.28.1 Detailed Description

Annotated Phylo Model.

A list of available terms for this model can be found in the Phylo counters section.

Class representing a phylogenetic tree model with annotations.

The Geese class represents a phylogenetic tree model with annotations. It includes a total of  $\mathbb{N}+1$  nodes, the +1 being the root node. The class provides methods for initializing the model, calculating the likelihood, simulating trees, and making predictions.

The class includes shared objects within a Geese object, such as rengine, model, states, n\_zeros, n\_ $\leftrightarrow$  ones, n\_dupl\_events, and n\_spec\_events. It also includes information about the type of event, such as etype\_default, etype\_speciation, etype\_duplication, and etype\_either.

The class provides constructors, a destructor, and methods for initializing the model, inheriting support, calculating the sequence, calculating the reduced sequence, calculating the likelihood, calculating the likelihood exhaustively, getting probabilities, setting the seed, simulating trees, parsing polytomies, getting observed counts, printing observed counts, printing information about the GEESE, and making predictions.

See also

Flock

Definition at line 114 of file geese-bones.hpp.

#### 8.28.2 Constructor & Destructor Documentation

#### 8.28.2.1 Geese() [1/4]

```
Geese::Geese ( ) [inline]
```

Definition at line 6 of file geese-meat-constructors.hpp.

## 8.28.2.2 Geese() [2/4]

```
Geese::Geese (
         std::vector< std::vector< size_t > & annotations,
         std::vector< size_t > & geneid,
         std::vector< int > & parent,
         std::vector< bool > & duplication ) [inline]
```

Definition at line 20 of file geese-meat-constructors.hpp.

## 8.28.2.3 Geese() [3/4]

Definition at line 230 of file geese-meat-constructors.hpp.

## 8.28.2.4 Geese() [4/4]

Definition at line 309 of file geese-meat-constructors.hpp.

#### 8.28.2.5 ∼Geese()

```
Geese::∼Geese ( ) [inline]
```

Definition at line 119 of file geese-meat.hpp.

## 8.28.3 Member Function Documentation

#### 8.28.3.1 calc\_reduced\_sequence()

```
void Geese::calc_reduced_sequence ( ) [inline]
```

Definition at line 361 of file geese-meat.hpp.

#### 8.28.3.2 calc\_sequence()

```
void Geese::calc_sequence (
          Node * n = nullptr ) [inline]
```

Definition at line 317 of file geese-meat.hpp.

## 8.28.3.3 colnames()

```
std::vector< std::string > Geese::colnames ( ) const [inline]
```

Names of the terms in the model.

Definition at line 483 of file geese-meat.hpp.

# 8.28.3.4 get\_annotated\_nodes()

```
std::vector< size_t > Geese::get_annotated_nodes ( ) const [inline]
```

Returns the ids of the nodes with at least one annotation.

Definition at line 772 of file geese-meat.hpp.

#### 8.28.3.5 get\_annotations()

```
std::vector< size_t > Geese::get_annotations ( ) const [inline]
```

Returns the annotations of the nodes with at least one annotation.

Definition at line 795 of file geese-meat.hpp.

#### 8.28.3.6 get\_counters()

```
PhyloCounters * Geese::get_counters ( ) [inline]
```

Definition at line 755 of file geese-meat.hpp.

#### 8.28.3.7 get\_model()

```
PhyloModel * Geese::get_model ( ) [inline]
```

Definition at line 760 of file geese-meat.hpp.

## 8.28.3.8 get\_probabilities()

```
std::vector< double > Geese::get_probabilities ( ) const [inline]
```

Definition at line 409 of file geese-meat.hpp.

# 8.28.3.9 get\_rengine()

```
std::mt19937 * Geese::get_rengine ( ) [inline]
```

Definition at line 750 of file geese-meat.hpp.

# 8.28.3.10 get\_states()

```
std::vector< std::vector< bool > > Geese::get_states ( ) const [inline]
```

Powerset of a gene's possible states.

This list of vectors is used throughout Geese. It lists all possible combinations of functional states for any gene. Thus, for P functions, there will be  $2^{P}$  possible combinations.

# Returns

```
std::vector< std::vector< bool > > of length 2^{^{\text{P}}}.
```

Definition at line 768 of file geese-meat.hpp.

8.28 Geese Class Reference 167

## 8.28.3.11 get\_support\_fun()

```
PhyloSupport * Geese::get_support_fun ( ) [inline]
```

Definition at line 764 of file geese-meat.hpp.

#### 8.28.3.12 inherit\_support()

Definition at line 260 of file geese-meat.hpp.

#### 8.28.3.13 init()

Definition at line 131 of file geese-meat.hpp.

#### 8.28.3.14 init\_node()

```
void Geese::init_node (
          Node & n ) [inline]
```

Definition at line 6 of file geese-meat.hpp.

#### 8.28.3.15 likelihood()

Definition at line 103 of file geese-meat-likelihood.hpp.

#### 8.28.3.16 likelihood\_exhaust()

Definition at line 7 of file geese-meat-likelihood\_exhaust.hpp.

## 8.28.3.17 nannotations()

```
std::vector< size_t > Geese::nannotations ( ) const [inline], [noexcept]
```

Number of annotations.

Definition at line 474 of file geese-meat.hpp.

## 8.28.3.18 nfuns()

```
size_t Geese::nfuns ( ) const [inline], [noexcept]
```

Number of functions analyzed.

Definition at line 430 of file geese-meat.hpp.

#### 8.28.3.19 nleafs()

```
size_t Geese::nleafs ( ) const [inline], [noexcept]
```

Number of leaf.

Definition at line 444 of file geese-meat.hpp.

## 8.28.3.20 nnodes()

```
size_t Geese::nnodes ( ) const [inline], [noexcept]
```

Number of nodes (interior + leaf)

Definition at line 437 of file geese-meat.hpp.

#### 8.28.3.21 nterms()

```
size_t Geese::nterms ( ) const [inline]
```

Number of terms included.

Definition at line 456 of file geese-meat.hpp.

## 8.28.3.22 observed\_counts()

```
std::vector< std::vector< double > > Geese::observed_counts ( ) [inline]
```

Definition at line 525 of file geese-meat.hpp.

#### 8.28.3.23 operator=() [1/2]

#### 8.28.3.24 operator=() [2/2]

```
Geese& Geese::operator= (
          Geese && model_ ) [delete], [noexcept]
```

## 8.28.3.25 parse\_polytomies()

Check polytomies and return the largest.

Definition at line 490 of file geese-meat.hpp.

#### 8.28.3.26 predict()

Definition at line 279 of file geese-meat-predict.hpp.

## 8.28.3.27 predict\_backend()

< True if the array belongs to the set

Definition at line 6 of file geese-meat-predict.hpp.

## 8.28.3.28 predict\_exhaust()

Definition at line 5 of file geese-meat-predict exhaust.hpp.

#### 8.28.3.29 predict\_exhaust\_backend()

Definition at line 47 of file geese-meat-predict exhaust.hpp.

#### 8.28.3.30 predict\_sim()

Definition at line 6 of file geese-meat-predict\_sim.hpp.

8.28 Geese Class Reference 171

#### 8.28.3.31 print()

```
void Geese::print ( ) const [inline]
```

Prints information about the GEESE.

Definition at line 659 of file geese-meat.hpp.

#### 8.28.3.32 print\_nodes()

```
void Geese::print_nodes ( ) const [inline]
```

Definition at line 677 of file geese-meat.hpp.

#### 8.28.3.33 print\_observed\_counts()

```
void Geese::print_observed_counts ( ) [inline]
```

Definition at line 596 of file geese-meat.hpp.

## 8.28.3.34 set\_seed()

Definition at line 4 of file geese-meat-simulate.hpp.

## 8.28.3.35 simulate()

```
\begin{tabular}{ll} {\tt std::vector}<& {\tt size\_t}>> {\tt Geese::simulate}~(\\ & {\tt const}~{\tt std::vector}<& {\tt double}>& {\tt par}~)~[{\tt inline}] \end{tabular}
```

Definition at line 8 of file geese-meat-simulate.hpp.

## 8.28.3.36 support\_size()

```
size_t Geese::support_size ( ) const [inline], [noexcept]
```

Number of unique sets of sufficient stats.

Definition at line 464 of file geese-meat.hpp.

#### 8.28.3.37 update\_annotations()

Definition at line 288 of file geese-meat.hpp.

#### 8.28.4 Member Data Documentation

#### 8.28.4.1 delete\_rengine

```
bool Geese::delete_rengine = false
```

Definition at line 153 of file geese-bones.hpp.

## 8.28.4.2 delete\_support

```
bool Geese::delete_support = false
```

Definition at line 154 of file geese-bones.hpp.

## 8.28.4.3 etype\_default

```
const size_t Geese::etype_default = 1ul [static]
```

Definition at line 167 of file geese-bones.hpp.

#### 8.28.4.4 etype\_duplication

```
const size_t Geese::etype_duplication = 1ul [static]
```

Definition at line 169 of file geese-bones.hpp.

## 8.28.4.5 etype\_either

```
const size_t Geese::etype_either = 2ul [static]
```

Definition at line 170 of file geese-bones.hpp.

#### 8.28.4.6 etype\_speciation

```
const size_t Geese::etype_speciation = Oul [static]
```

Definition at line 168 of file geese-bones.hpp.

#### 8.28.4.7 initialized

```
bool Geese::initialized = false
```

Definition at line 152 of file geese-bones.hpp.

# 8.28.4.8 map\_to\_state\_id

```
barry::MapVec_type< size_t > Geese::map_to_state_id
```

Definition at line 144 of file geese-bones.hpp.

#### 8.28.4.9 nfunctions

```
size_t Geese::nfunctions
```

Definition at line 141 of file geese-bones.hpp.

#### 8.28.4.10 nodes

```
std::map< size_t, Node > Geese::nodes
```

Definition at line 142 of file geese-bones.hpp.

#### 8.28.4.11 pset\_loc

```
std::vector< std::vector< size_t > > > Geese::pset_loc
```

Locations of columns.

Definition at line 145 of file geese-bones.hpp.

#### 8.28.4.12 reduced\_sequence

```
std::vector< size_t > Geese::reduced_sequence
```

Definition at line 149 of file geese-bones.hpp.

#### 8.28.4.13 sequence

```
std::vector< size_t > Geese::sequence
```

Definition at line 148 of file geese-bones.hpp.

The documentation for this class was generated from the following files:

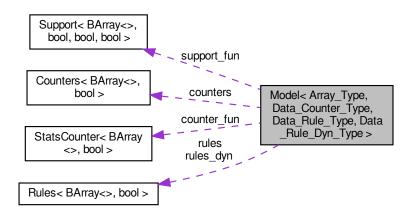
- include/barry/models/geese/geese-bones.hpp
- include/barry/models/geese/geese-meat-constructors.hpp
- include/barry/models/geese/geese-meat-likelihood.hpp
- include/barry/models/geese/geese-meat-likelihood\_exhaust.hpp
- include/barry/models/geese/geese-meat-predict.hpp
- include/barry/models/geese/geese-meat-predict\_exhaust.hpp
- include/barry/models/geese/geese-meat-predict\_sim.hpp
- include/barry/models/geese/geese-meat-simulate.hpp
- include/barry/models/geese/geese-meat.hpp

# 8.29 Model < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type > Class Template Reference

General framework for discrete exponential models. This class allows generating discrete exponential models in the form of a linear exponential model:

#include <model-bones.hpp>

Collaboration diagram for Model < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type >:



#### **Public Member Functions**

- void update\_normalizing\_constants (const std::vector< double > &params,)
  - Computes the normalizing constant for a given set of parameters.
- void  $update\_likelihoods$  (const std::vector < double > &params,)
- void update pset probs (const std::vector< double > &params,)
- void set rengine (std::mt19937 \*rengine , bool delete =false)
- void set\_seed (size\_t s)
- Model ()
- Model (size\_t size\_)
- Model (const Model < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type > &Model ← \_\_)
- Model< Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type > & operator= (const Model< Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type > &Model\_)
- virtual ∼Model ()
- void store\_psets () noexcept
- std::vector< double > gen\_key (const Array\_Type &Array\_)
- size\_t add\_array (const Array\_Type &Array\_, bool force\_new=false)

Adds an array to the support of not already included.

- · void print\_stats (size\_t i) const
- virtual void print () const

Prints information about the model.

- Array\_Type sample (const Array\_Type &Array\_, const std::vector< double > &params={})
- Array\_Type sample (const size\_t &i, const std::vector< double > &params)

double conditional\_prob (const Array\_Type &Array\_, const std::vector< double > &params, size\_t i, size\_t j)
 Conditional probability ("Gibbs sampler")

- const std::mt19937 \* get\_rengine () const
- Counters < Array Type, Data Counter Type > \* get counters ()
- Rules
   Array\_Type, Data\_Rule\_Type > \* get\_rules ()
- Rules< Array\_Type, Data\_Rule\_Dyn\_Type > \* get\_rules\_dyn ()
- Support< Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type > \* get\_support\_fun ()

#### Wrappers for the <tt>Counters</tt> member.

These will add counters to the model, which are shared by the support and the actual counter function.

- void add counter (Counter< Array Type, Data Counter Type > &counter)
- void add\_counter (Counter\_fun\_type < Array\_Type, Data\_Counter\_Type > count\_fun\_, Counter\_fun\_type 
   Array\_Type, Data\_Counter\_Type > init\_fun\_=nullptr, Data\_Counter\_Type data\_=nullptr)
- void set\_counters (Counters < Array\_Type, Data\_Counter\_Type > \*counters\_)
- void add hasher (Hasher fun type< Array Type, Data Counter Type > fun )

#### Wrappers for the <tt>Rules</tt> member.

These will add rules to the model, which are shared by the support and the actual counter function.

- void add\_rule (Rule < Array\_Type, Data\_Rule\_Type > &rule)
- void add\_rule (Rule\_fun\_type < Array\_Type, Data\_Rule\_Type > count\_fun\_, Data\_Rule\_Type data\_)
- void set\_rules (Rules < Array\_Type, Data\_Rule\_Type > \*rules\_)
- void add\_rule\_dyn (Rule< Array\_Type, Data\_Rule\_Dyn\_Type > &rule)
- void set\_rules\_dyn (Rules < Array\_Type, Data\_Rule\_Dyn\_Type > \*rules\_)

#### Likelihood functions.

Calculation of likelihood functions is done reusing normalizing constants. Before recalculating the normalizing constant, the function checks whether params matches the last set vector of parameters used to compute it.

#### Parameters

params	Vector of parameters
as_log	When true, the function returns the log-likelihood.

- double likelihood (const std::vector< double > &params, const size\_t &i, bool as\_log=false, bool no\_

   update normconst=false)
- double likelihood (const std::vector< double > &params, const std::vector< double > &target\_, const size t &i, bool as log=false, bool no update normconst=false)
- double likelihood (const std::vector< double > &params, const double \*target\_, const size\_t &i, bool as 
   log=false, bool no update normconst=false)
- double likelihood\_total (const std::vector< double > &params, bool as\_log=false, BARRY\_NCORES\_ARG(=2), bool no\_update\_normconst=false)

#### Extract elements by index

#### Parameters

i	Index relative to the array in the model.
params	A new vector of model parameters to compute the normalizing constant.
as_log	When true returns the logged version of the normalizing constant.

```
• const std::vector< double > & get_normalizing_constants () const
```

- const std::vector< double > & get\_likelihoods () const
- const std::vector< Array Type > \* get pset (const size t &i)
- const double \* get\_pset\_stats (const size\_t &i)

#### Size of the model

Number of different supports included in the model

This will return the size of stats\_target.

#### Returns

size() returns the number of arrays in the model.
size\_unique() returns the number of unique arrays (according to the hasher) in the model.
nterms() returns the number of terms in the model.

- size\_t size () const noexcept
- size\_t size\_unique () const noexcept
- size\_t nterms () const noexcept
- size\_t nrules () const noexcept
- size\_t nrules\_dyn () const noexcept
- size t support size () const noexcept
- std::vector< std::string > colnames () const

```
    std::vector< std::vector< double > > * get stats target ()
```

Raw pointers to the support and target statistics.

std::vector< double > \* get stats support ()

Sufficient statistics of the support(s)

std::vector< size\_t > \* get\_stats\_support\_sizes ()

Number of vectors included in the support.

std::vector< size\_t > \* get\_stats\_support\_sizes\_acc ()

Accumulated number of vectors included in the support.

- $std::vector < size_t > * get_arrays2support ()$
- std::vector< std::vector< Array\_Type > > \* get\_pset\_arrays ()
- std::vector< double > \* get\_pset\_stats ()

Statistics of the support(s)

- std::vector< double > \* get\_pset\_probs ()
- std::vector< size\_t > \* get\_pset\_sizes ()
- std::vector< size\_t > \* get\_pset\_locations ()

void set\_transform\_model (std::function < std::vector < double > (double \*, size\_t) > fun, std::vector < std ← ::string > names)

Set the transform\_model\_fun object.

std::vector< double > transform\_model (double \*data, size\_t k)

#### **Protected Attributes**

MapVec\_type< double, size\_t > keys2support

Map of types of arrays to support sets.

std::vector< std::vector< double >> params last

Vector of the previously used parameters.

- std::vector< double > normalizing\_constants
- std::vector< bool > first\_calc\_done
- bool delete counters = false
- bool delete rules = false
- bool delete rules dyn = false
- $\bullet \ \ \, std:: function < std:: vector < double > double *, size\_t \ k) > transform\_model\_fun = nullptr$

Transformation of the model.

std::vector< std::string > transform model term names

#### Random number generation

Random number generation

- std::mt19937 \* rengine = nullptr
- bool delete rengine = false

#### Information about the arrays used in the model

stats\_target holds the observed sufficient statistics for each array in the dataset. array\_← frequency contains the frequency with which each of the target stats\_target (arrays) shows in the support. array2support maps array indices (0, 1, ...) to the corresponding support.

Each vector of stats\_support has the data stored in a row-wise order, with each row starting with the weights, e.g., in a model with k terms the first k + 1 elements of stats\_support would be:

- · weights
- term 1
- term 2
- ...
- term k
- std::vector< double > stats support

Sufficient statistics of the model (support)

 $\bullet \ \ \mathsf{std} :: \mathsf{vector} < \mathsf{size\_t} > \mathsf{stats\_support\_sizes}$ 

Number of vectors included in the support.

• std::vector< size\_t > stats\_support\_sizes\_acc

Accumulated number of vectors included in the support.

std::vector< size\_t > stats\_support\_n\_arrays

Number of arrays included per support.

std::vector< std::vector< double >> stats\_target

Target statistics of the model.

- std::vector< double > stats\_likelihood
- std::vector< size\_t > arrays2support

## Container space for the powerset (and its sufficient stats\_target)

This is useful in the case of using simulations or evaluating functions that need to account for the full set of states.

```
• bool with_pset = false
```

std::vector< std::vector< Array\_Type >> pset\_arrays
 Arrays of the support(s)

 $\bullet \ \, \mathsf{std} : \!\! \mathsf{vector} \!\! < \mathsf{double} > \mathsf{pset\_stats}$ 

Statistics of the support(s)

std::vector< double > pset\_probs

Probabilities of the support(s)

• std::vector< size\_t > pset\_sizes

Number of vectors included in the support.

• std::vector< size t > pset locations

Accumulated number of vectors included in the support.

#### **Functions to compute statistics**

Arguments are recycled to save memory and computation.

- Counters
   Array\_Type, Data\_Counter\_Type > \* counters
- Rules < Array\_Type, Data\_Rule\_Type > \* rules
- Rules< Array\_Type, Data\_Rule\_Dyn\_Type > \* rules\_dyn
- Support < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type > support\_fun
- StatsCounter< Array\_Type, Data\_Counter\_Type > counter\_fun

# 8.29.1 Detailed Description

template<typename Array\_Type = BArray<>>, typename Data\_Counter\_Type = bool, typename Data\_Rule\_Type = bool, typename Data\_Rule\_Dyn\_Type = bool>

class Model < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type >

General framework for discrete exponential models. This class allows generating discrete exponential models in the form of a linear exponential model:

$$\frac{\exp\left(\theta^{\mathsf{t}}c(A)\right)}{\sum_{A'\in\mathcal{A}}\exp\left(\theta^{\mathsf{t}}c(A')\right)}$$

This implementation aims to reduce the number of times that the support needs to be computed. Models included here use more than a single array, and thus allow the function to recycle support sets as needed. For example, if we are looking at directed graphs all of the same size and without vertex level features, i.e. a model that only counts edges, triangles, etc. then the support needs to be fully computed only once.

#### **Template Parameters**

Array_Type	Class of BArray object.
Data_Counter_Type	Any type.
Data_Rule_Type	Any type.

Definition at line 34 of file model-bones.hpp.

#### 8.29.2 Constructor & Destructor Documentation

#### 8.29.2.1 Model() [1/3]

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::Model [inline]
```

Definition at line 288 of file model-meat.hpp.

#### 8.29.2.2 Model() [2/3]

Definition at line 326 of file model-meat.hpp.

# 8.29.2.3 Model() [3/3]

Definition at line 368 of file model-meat.hpp.

#### 8.29.2.4 ∼Model()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
virtual Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::~Model (
) [inline], [virtual]
```

Definition at line 185 of file model-bones.hpp.

# 8.29.3 Member Function Documentation

## 8.29.3.1 add\_array()

Adds an array to the support of not already included.

#### **Parameters**

Array_	array to be added
force_new	If false, it will use keygen to obtain a double vector and create a hash of it. If the hash has
	been computed earlier, the support is recycled.

#### Returns

The number of the array.

When computing with the powerset, we need to grow the corresponding vectors on the fly

Definition at line 612 of file model-meat.hpp.

#### 8.29.3.2 add\_counter() [1/2]

Definition at line 487 of file model-meat.hpp.

## 8.29.3.3 add\_counter() [2/2]

Definition at line 496 of file model-meat.hpp.

## 8.29.3.4 add\_hasher()

Definition at line 531 of file model-meat.hpp.

## 8.29.3.5 add\_rule() [1/2]

Definition at line 542 of file model-meat.hpp.

#### 8.29.3.6 add rule() [2/2]

#### 8.29.3.7 add rule dyn() [1/2]

Definition at line 570 of file model-meat.hpp.

#### 8.29.3.8 add rule dyn() [2/2]

Definition at line 579 of file model-meat.hpp.

#### 8.29.3.9 colnames()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< std::string > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Cyn_Type >::colnames [inline]
```

Definition at line 1350 of file model-meat.hpp.

## 8.29.3.10 conditional\_prob()

Conditional probability ("Gibbs sampler")

Computes the conditional probability of observing  $P\{Y(i,j) = | Y^C, \text{ theta}\}$ , i.e., the probability of observing the entry Y(i,j) equal to one given the rest of the array.

#### **Parameters**

Array⊷	Array to check
_	
params	Vector of parameters
i	Row entry
j	Column entry

#### Returns

double The conditional probability

Definition at line 1646 of file model-meat.hpp.

#### 8.29.3.11 gen key()

Definition at line 480 of file model-meat.hpp.

## 8.29.3.12 get\_arrays2support()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< size_t > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn
_Type >::get_arrays2support [inline]
```

Definition at line 1735 of file model-meat.hpp.

## 8.29.3.13 get\_counters()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
Counters< Array_Type, Data_Counter_Type > * Model< Array_Type, Data_Counter_Type, Data_Rule
_Type, Data_Rule_Dyn_Type >::get_counters [inline]
```

Definition at line 1684 of file model-meat.hpp.

#### 8.29.3.14 get\_likelihoods()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
const std::vector< double > & Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_←
Rule_Dyn_Type >::get_likelihoods [inline]
```

Definition at line 1157 of file model-meat.hpp.

#### 8.29.3.15 get\_normalizing\_constants()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
const std::vector< double > & Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_\top Rule_Dyn_Type >::get_normalizing_constants [inline]
```

Definition at line 1144 of file model-meat.hpp.

#### 8.29.3.16 get\_pset()

Definition at line 1165 of file model-meat.hpp.

## 8.29.3.17 get\_pset\_arrays()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< std::vector< Array_Type > > * Model< Array_Type, Data_Counter_Type, Data_Rule_\Lorenty
Type, Data_Rule_Dyn_Type >::get_pset_arrays [inline]
```

Definition at line 1742 of file model-meat.hpp.

## 8.29.3.18 get\_pset\_locations()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< size_t > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn 
_Type >::get_pset_locations [inline]
```

Definition at line 1780 of file model-meat.hpp.

## 8.29.3.19 get\_pset\_probs()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< double > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn 
_Type >::get_pset_probs [inline]
```

Definition at line 1759 of file model-meat.hpp.

#### 8.29.3.20 get pset sizes()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< size_t > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn
_Type >::get_pset_sizes [inline]
```

Definition at line 1769 of file model-meat.hpp.

## 8.29.3.21 get\_pset\_stats() [1/2]

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< double > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn 
_Type >::get_pset_stats [inline]
```

Statistics of the support(s)

Definition at line 1748 of file model-meat.hpp.

#### 8.29.3.22 get\_pset\_stats() [2/2]

Definition at line 1179 of file model-meat.hpp.

#### 8.29.3.23 get\_rengine()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
const std::mt19937 * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type
>::get_rengine [inline]
```

Definition at line 1679 of file model-meat.hpp.

## 8.29.3.24 get\_rules()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
Rules< Array_Type, Data_Rule_Type > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_rules [inline]
```

Definition at line 1689 of file model-meat.hpp.

#### 8.29.3.25 get\_rules\_dyn()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >

Rules< Array_Type, Data_Rule_Dyn_Type > * Model< Array_Type, Data_Counter_Type, Data_Rule_\top Type, Data_Rule_Dyn_Type >::get_rules_dyn [inline]
```

Definition at line 1694 of file model-meat.hpp.

#### 8.29.3.26 get\_stats\_support()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< double > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn 
_Type >::get_stats_support [inline]
```

Sufficient statistics of the support(s)

Definition at line 1712 of file model-meat.hpp.

#### 8.29.3.27 get\_stats\_support\_sizes()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< size_t > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn 
_Type >::get_stats_support_sizes [inline]
```

Number of vectors included in the support.

Definition at line 1720 of file model-meat.hpp.

### 8.29.3.28 get\_stats\_support\_sizes\_acc()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< size_t > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn
_Type >::get_stats_support_sizes_acc [inline]
```

Accumulated number of vectors included in the support.

Definition at line 1728 of file model-meat.hpp.

#### 8.29.3.29 get\_stats\_target()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
std::vector< std::vector< double > > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_stats_target [inline]
```

Raw pointers to the support and target statistics.

The support of the model is stored as a vector of vector < double>. Each element of it contains the support for an specific type of array included. It represents an array of size  $(k + 1) \times n \text{ unique elements}$ , with the data stored by-row. The last element of each entry corresponds to the weights, i.e., the frequency with which such sufficient statistics are observed in the support.

Definition at line 1705 of file model-meat.hpp.

## 8.29.3.30 get\_support\_fun()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > * Model< Array

_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::get_support_fun [inline]
```

Definition at line 1700 of file model-meat.hpp.

## 8.29.3.31 likelihood() [1/4]

Definition at line 836 of file model-meat.hpp.

#### 8.29.3.32 likelihood() [2/4]

Definition at line 986 of file model-meat.hpp.

## 8.29.3.33 likelihood() [3/4]

Definition at line 789 of file model-meat.hpp.

#### 8.29.3.34 likelihood() [4/4]

Definition at line 920 of file model-meat.hpp.

#### 8.29.3.35 likelihood total()

Definition at line 1058 of file model-meat.hpp.

#### 8.29.3.36 nrules()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
size_t Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::nrules
[inline], [noexcept]
```

Definition at line 1320 of file model-meat.hpp.

## 8.29.3.37 nrules\_dyn()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
size_t Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::nrules_dyn
[inline], [noexcept]
```

Definition at line 1328 of file model-meat.hpp.

#### 8.29.3.38 nterms()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
size_t Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::nterms
[inline], [noexcept]
```

Definition at line 1309 of file model-meat.hpp.

## 8.29.3.39 operator=()

Definition at line 418 of file model-meat.hpp.

## 8.29.3.40 print()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::print [inline],
[virtual]
```

Prints information about the model.

Definition at line 1231 of file model-meat.hpp.

#### 8.29.3.41 print\_stats()

Definition at line 1191 of file model-meat.hpp.

## 8.29.3.42 sample() [1/2]

When computing with the powerset, we need to grow the corresponding vectors on the fly

Definition at line 1433 of file model-meat.hpp.

#### 8.29.3.43 sample() [2/2]

Definition at line 1367 of file model-meat.hpp.

## 8.29.3.44 set\_counters()

Definition at line 513 of file model-meat.hpp.

#### 8.29.3.45 set\_rengine()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::set_rengine (
    std::mt19937 * rengine_,
    bool delete_ = false ) [inline]
```

Definition at line 155 of file model-bones.hpp.

## 8.29.3.46 set\_rules()

Definition at line 552 of file model-meat.hpp.

## 8.29.3.47 set\_rules\_dyn()

Definition at line 594 of file model-meat.hpp.

#### 8.29.3.48 set\_seed()

Definition at line 165 of file model-bones.hpp.

#### 8.29.3.49 set\_transform\_model()

Set the transform\_model\_fun object.

The transform\_model function is used to transform the data

#### **Parameters**

data	
target	
n_arrays	
arrays2support	

Definition at line 1792 of file model-meat.hpp.

## 8.29.3.50 size()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
size_t Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::size [inline],
[noexcept]
```

Definition at line 1292 of file model-meat.hpp.

#### 8.29.3.51 size\_unique()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
size_t Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::size_unique
[inline], [noexcept]
```

Definition at line 1300 of file model-meat.hpp.

#### 8.29.3.52 store\_psets()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::store_psets
[inline], [noexcept]
```

Definition at line 474 of file model-meat.hpp.

#### 8.29.3.53 support\_size()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
size_t Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::support_
size [inline], [noexcept]
```

Definition at line 1336 of file model-meat.hpp.

#### 8.29.3.54 transform\_model()

### 8.29.3.55 update\_likelihoods()

Definition at line 176 of file model-meat.hpp.

### 8.29.3.56 update\_normalizing\_constants()

Computes the normalizing constant for a given set of parameters.

This function will compute the normalizing constant for a given set of parameters. It will also update the normalizing\_constants member variable.

Definition at line 134 of file model-meat.hpp.

## 8.29.3.57 update\_pset\_probs()

Definition at line 215 of file model-meat.hpp.

#### 8.29.4 Member Data Documentation

#### 8.29.4.1 arrays2support

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< size_t > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::arrays2support [protected]
```

Definition at line 68 of file model-bones.hpp.

#### 8.29.4.2 counter fun

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
StatsCounter<Array_Type, Data_Counter_Type> Model< Array_Type, Data_Counter_Type, Data_Rule_←
Type, Data_Rule_Dyn_Type >::counter_fun [protected]
```

Definition at line 100 of file model-bones.hpp.

#### 8.29.4.3 counters

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Counters<Array_Type, Data_Counter_Type>* Model< Array_Type, Data_Counter_Type, Data_Rule_Dyn_Type >::counters [protected]
```

Definition at line 96 of file model-bones.hpp.

## 8.29.4.4 delete\_counters

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_←
counters = false [protected]
```

Definition at line 108 of file model-bones.hpp.

#### 8.29.4.5 delete\_rengine

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_←
rengine = false [protected]
```

Definition at line 43 of file model-bones.hpp.

#### 8.29.4.6 delete rules

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_rules
= false [protected]
```

Definition at line 109 of file model-bones.hpp.

#### 8.29.4.7 delete rules dyn

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_←
rules_dyn = false [protected]
```

Definition at line 110 of file model-bones.hpp.

## 8.29.4.8 first\_calc\_done

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< bool > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type
>::first_calc_done [protected]
```

Definition at line 106 of file model-bones.hpp.

#### 8.29.4.9 keys2support

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
MapVec_type< double, size_t > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_\times
Rule_Dyn_Type >::keys2support [protected]
```

Map of types of arrays to support sets.

This is of the same length as the vector stats\_target.

Definition at line 75 of file model-bones.hpp.

## 8.29.4.10 normalizing\_constants

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< double > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::normalizing_constants [protected]
```

Definition at line 105 of file model-bones.hpp.

## 8.29.4.11 params\_last

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector<double> > Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::params_last [protected]
```

Vector of the previously used parameters.

Definition at line 104 of file model-bones.hpp.

#### 8.29.4.12 pset\_arrays

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector< Array_Type >> Model< Array_Type, Data_Counter_Type, Data_Rule_←
Type, Data_Rule_Dyn_Type >::pset_arrays [protected]
```

Arrays of the support(s)

Definition at line 84 of file model-bones.hpp.

#### 8.29.4.13 pset locations

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< size_t > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::pset_locations [protected]
```

Accumulated number of vectors included in the support.

Definition at line 88 of file model-bones.hpp.

#### 8.29.4.14 pset\_probs

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< double > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::pset_probs [protected]
```

Probabilities of the support(s)

Definition at line 86 of file model-bones.hpp.

#### 8.29.4.15 pset sizes

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< size_t > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::pset_sizes [protected]
```

Number of vectors included in the support.

Definition at line 87 of file model-bones.hpp.

## 8.29.4.16 pset\_stats

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< double > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::pset_stats [protected]
```

Statistics of the support(s)

Definition at line 85 of file model-bones.hpp.

#### 8.29.4.17 rengine

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::mt19937* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >←
::rengine = nullptr [protected]
```

Definition at line 42 of file model-bones.hpp.

#### 8.29.4.18 rules

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Rules<Array_Type, Data_Rule_Type>* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data←
_Rule_Dyn_Type >::rules [protected]
```

Definition at line 97 of file model-bones.hpp.

## 8.29.4.19 rules\_dyn

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Rules<Array_Type, Data_Rule_Dyn_Type>* Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::rules_dyn [protected]
```

Definition at line 98 of file model-bones.hpp.

#### 8.29.4.20 stats likelihood

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< double > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::stats_likelihood [protected]
```

Definition at line 67 of file model-bones.hpp.

## 8.29.4.21 stats\_support

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< double > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::stats_support [protected]
```

Sufficient statistics of the model (support)

Definition at line 62 of file model-bones.hpp.

#### 8.29.4.22 stats\_support\_n\_arrays

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< size_t > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::stats_support_n_arrays [protected]
```

Number of arrays included per support.

Definition at line 65 of file model-bones.hpp.

#### 8.29.4.23 stats support sizes

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< size_t > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::stats_support_sizes [protected]
```

Number of vectors included in the support.

Definition at line 63 of file model-bones.hpp.

## 8.29.4.24 stats\_support\_sizes\_acc

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< size_t > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::stats_support_sizes_acc [protected]
```

Accumulated number of vectors included in the support.

Definition at line 64 of file model-bones.hpp.

#### 8.29.4.25 stats\_target

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector< double > > Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::stats_target [protected]
```

Target statistics of the model.

Definition at line 66 of file model-bones.hpp.

## 8.29.4.26 support\_fun

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Support<Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type> Model< Array_Type,
Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::support_fun [protected]
```

Definition at line 99 of file model-bones.hpp.

## 8.29.4.27 transform\_model\_fun

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::function<std::vector<double>double *, size_t k)> Model< Array_Type, Data_Counter_Type,
Data_Rule_Type, Data_Rule_Dyn_Type >::transform_model_fun = nullptr [protected]
```

Transformation of the model.

When specified, this function will update the model by modifying the linear equation. For example, if the user wanted to add interaction terms, rescale, or apply other operations of the sorts, the user can do such through this function.

The function should return void and receive the following arguments:

- data Pointer to the first element of the set of sufficient statistics
- · k size\_t indicating the number of sufficient statistics

#### Returns

Nothing, but it will modify the model data.

Definition at line 128 of file model-bones.hpp.

## 8.29.4.28 transform\_model\_term\_names

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::string > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_\times
Dyn_Type >::transform_model_term_names [protected]
```

Definition at line 130 of file model-bones.hpp.

## 8.29.4.29 with\_pset

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::with_pset =
false [protected]
```

Definition at line 83 of file model-bones.hpp.

The documentation for this class was generated from the following files:

- include/barry/model-bones.hpp
- include/barry/model-meat.hpp

## 8.30 NetCounterData Class Reference

Data class used to store arbitrary size\_t or double vectors.

```
#include <network.hpp>
```

#### **Public Member Functions**

- · NetCounterData ()
- NetCounterData (const std::vector < size\_t > &indices\_, const std::vector < double > &numbers\_)
- ∼NetCounterData ()

## **Public Attributes**

- std::vector< size\_t > indices
- std::vector< double > numbers

## 8.30.1 Detailed Description

Data class used to store arbitrary size\_t or double vectors.

Definition at line 56 of file network.hpp.

## 8.30.2 Constructor & Destructor Documentation

## 8.30.2.1 NetCounterData() [1/2]

```
NetCounterData::NetCounterData ( ) [inline]
```

Definition at line 62 of file network.hpp.

## 8.30.2.2 NetCounterData() [2/2]

Definition at line 63 of file network.hpp.

## 8.30.2.3 ~NetCounterData()

```
NetCounterData::~NetCounterData ( ) [inline]
```

Definition at line 68 of file network.hpp.

#### 8.30.3 Member Data Documentation

#### 8.30.3.1 indices

```
std::vector< size_t > NetCounterData::indices
```

Definition at line 59 of file network.hpp.

#### 8.30.3.2 numbers

```
std::vector< double > NetCounterData::numbers
```

Definition at line 60 of file network.hpp.

The documentation for this class was generated from the following file:

• include/barry/counters/network.hpp

## 8.31 NetworkData Class Reference

Data class for Networks.

```
#include <network.hpp>
```

#### **Public Member Functions**

- · NetworkData ()
- NetworkData (std::vector< double > vertex\_attr\_, bool directed\_=true)

Constructor using a single attribute.

NetworkData (std::vector< std::vector< double >> vertex\_attr\_, bool directed\_=true)

Constructor using multiple attributes.

∼NetworkData ()

#### **Public Attributes**

- bool directed = true
- std::vector< std::vector< double >> vertex attr

## 8.31.1 Detailed Description

Data class for Networks.

Details on the available counters for NetworkData can be found in the Network counters section.

This holds information about whether the graph is directed or not, and, if defined, vectors of node (vertex) attributes (vertex\_attr).

Definition at line 19 of file network.hpp.

## 8.31.2 Constructor & Destructor Documentation

#### 8.31.2.1 NetworkData() [1/3]

```
NetworkData::NetworkData ( ) [inline]
```

Definition at line 25 of file network.hpp.

## 8.31.2.2 NetworkData() [2/3]

Constructor using a single attribute.

#### **Parameters**

vertex_←	Double vector of length equal to the number of vertices in the data.
attr_	
directed_	When true the graph as treated as directed.

Definition at line 33 of file network.hpp.

## 8.31.2.3 NetworkData() [3/3]

```
NetworkData::NetworkData (
          std::vector< std::vector< double > > vertex_attr_,
          bool directed_ = true ) [inline]
```

Constructor using multiple attributes.

#### **Parameters**

vertex_←	Vector of double vectors. The size equals to the number of attributes to be created. Each
attr_	individual vector should be of length equal to the number of vertices.
directed_	When true the graph as treated as directed.

Definition at line 45 of file network.hpp.

## 8.31.2.4 ~NetworkData()

```
NetworkData::~NetworkData ( ) [inline]
```

Definition at line 51 of file network.hpp.

## 8.31.3 Member Data Documentation

## 8.31.3.1 directed

```
bool NetworkData::directed = true
```

Definition at line 22 of file network.hpp.

## 8.31.3.2 vertex\_attr

```
std::vector< std::vector< double > > NetworkData::vertex_attr
```

Definition at line 23 of file network.hpp.

The documentation for this class was generated from the following file:

• include/barry/counters/network.hpp

## 8.32 Node Class Reference

A single node for the model.

```
#include <geese-node-bones.hpp>
```

Collaboration diagram for Node:



## **Public Member Functions**

- ∼Node ()
- int get\_parent () const
- size\_t noffspring () const noexcept
- bool is\_leaf () const noexcept

## Construct a new Node object

- Node ()
- Node (size\_t id\_, size\_t ord\_, bool duplication\_)
- Node (size\_t id\_, size\_t ord\_, std::vector < size\_t > annotations\_, bool duplication\_)
- Node (Node &&x) noexcept
- Node (const Node &x)

8.32 Node Class Reference 207

#### **Public Attributes**

```
size_t id
```

Id of the node (as specified in the input)

· size tord

Order in which the node was created.

PhyloArray array

Array of the node.

std::vector< size\_t > annotations

Observed annotations (only defined for Geese)

- bool duplication
- std::vector< PhyloArray > arrays = {}

Arrays given all possible states.

Node \* parent = nullptr

Parent node.

std::vector< Node \* > offspring = {}

Offspring nodes.

• std::vector< size\_t > narray = {}

ID of the array in the model.

- bool visited = false
- std::vector< double > subtree\_prob

Induced subtree probabilities.

std::vector< double > probability

The probability of observing each state.

## 8.32.1 Detailed Description

A single node for the model.

Each node contains all the information to compute the conditional probability of the pruning algorithm at that node.

Definition at line 11 of file geese-node-bones.hpp.

#### 8.32.2 Constructor & Destructor Documentation

#### 8.32.2.1 Node() [1/5]

```
Node::Node ( ) [inline]
```

Definition at line 37 of file geese-node-bones.hpp.

## 8.32.2.2 Node() [2/5]

Definition at line 57 of file geese-node-bones.hpp.

## 8.32.2.3 Node() [3/5]

Definition at line 63 of file geese-node-bones.hpp.

## 8.32.2.4 Node() [4/5]

Definition at line 70 of file geese-node-bones.hpp.

## 8.32.2.5 Node() [5/5]

Definition at line 85 of file geese-node-bones.hpp.

## 8.32.2.6 ∼Node()

```
Node::~Node ( ) [inline]
```

Definition at line 48 of file geese-node-bones.hpp.

## 8.32.3 Member Function Documentation

8.32 Node Class Reference 209

## 8.32.3.1 get\_parent()

```
int Node::get_parent ( ) const [inline]
```

Definition at line 100 of file geese-node-bones.hpp.

## 8.32.3.2 is\_leaf()

```
bool Node::is_leaf ( ) const [inline], [noexcept]
```

Definition at line 112 of file geese-node-bones.hpp.

## 8.32.3.3 noffspring()

```
size_t Node::noffspring ( ) const [inline], [noexcept]
```

Definition at line 106 of file geese-node-bones.hpp.

### 8.32.4 Member Data Documentation

## 8.32.4.1 annotations

```
std::vector< size_t > Node::annotations
```

Observed annotations (only defined for Geese)

Definition at line 18 of file geese-node-bones.hpp.

#### 8.32.4.2 array

PhyloArray Node::array

Array of the node.

Definition at line 17 of file geese-node-bones.hpp.

## 8.32.4.3 arrays

```
std::vector< PhyloArray > Node::arrays = {}
```

Arrays given all possible states.

Definition at line 21 of file geese-node-bones.hpp.

## 8.32.4.4 duplication

```
bool Node::duplication
```

Definition at line 19 of file geese-node-bones.hpp.

## 8.32.4.5 id

```
size_t Node::id
```

Id of the node (as specified in the input)

Definition at line 14 of file geese-node-bones.hpp.

## 8.32.4.6 narray

```
std::vector< size_t > Node::narray = {}
```

ID of the array in the model.

Definition at line 25 of file geese-node-bones.hpp.

#### 8.32.4.7 offspring

```
std::vector< Node* > Node::offspring = {}
```

Offspring nodes.

Definition at line 24 of file geese-node-bones.hpp.

8.32 Node Class Reference 211

#### 8.32.4.8 ord

```
size_t Node::ord
```

Order in which the node was created.

Definition at line 15 of file geese-node-bones.hpp.

#### 8.32.4.9 parent

```
Node* Node::parent = nullptr
```

Parent node.

Definition at line 23 of file geese-node-bones.hpp.

## 8.32.4.10 probability

```
std::vector< double > Node::probability
```

The probability of observing each state.

Definition at line 29 of file geese-node-bones.hpp.

## 8.32.4.11 subtree\_prob

```
std::vector< double > Node::subtree_prob
```

Induced subtree probabilities.

Definition at line 28 of file geese-node-bones.hpp.

#### 8.32.4.12 visited

```
bool Node::visited = false
```

Definition at line 26 of file geese-node-bones.hpp.

The documentation for this class was generated from the following file:

include/barry/models/geese/geese-node-bones.hpp

## 8.33 NodeData Class Reference

Data definition for the PhyloArray class.

```
#include <geese-types.hpp>
```

## **Public Member Functions**

NodeData (const std::vector< double > &blengths\_, const std::vector< bool > &states\_, bool duplication
 —=true, bool has\_leaf\_=false)

## **Public Attributes**

```
    std::vector< double > blengths = {}
    std::vector< bool > states = {}
    bool duplication = true
        Whether the node is a duplication.
    bool has_leaf = false
        Whether the node has a leaf as offspring.
```

## 8.33.1 Detailed Description

Data definition for the PhyloArray class.

This holds basic information about a given node.

Definition at line 15 of file geese-types.hpp.

## 8.33.2 Constructor & Destructor Documentation

## 8.33.2.1 NodeData()

Definition at line 32 of file geese-types.hpp.

## 8.33.3 Member Data Documentation

## 8.33.3.1 blengths

```
std::vector< double > NodeData::blengths = {}
```

Branch length.

Definition at line 21 of file geese-types.hpp.

## 8.33.3.2 duplication

```
bool NodeData::duplication = true
```

Whether the node is a duplication.

Definition at line 28 of file geese-types.hpp.

## 8.33.3.3 has\_leaf

```
bool NodeData::has_leaf = false
```

Whether the node has a leaf as offspring.

Definition at line 29 of file geese-types.hpp.

#### 8.33.3.4 states

```
std::vector< bool > NodeData::states = {}
```

State of the parent node.

Definition at line 26 of file geese-types.hpp.

The documentation for this class was generated from the following file:

• include/barry/models/geese/geese-types.hpp

# 8.34 PhyloCounterData Class Reference

```
#include <geese-types.hpp>
```

#### **Public Member Functions**

```
PhyloCounterData (std::vector< size_t > data_, std::vector< double > *counters_=nullptr)
PhyloCounterData ()
size_t at (size_t d)
size_t operator() (size_t d)
size_t operator[] (size_t d)
void reserve (size_t x)
void push_back (size_t x)
void shrink_to_fit ()
size_t size ()
std::vector< size_t >::iterator begin ()
std::vector< size_t >::iterator end ()
bool empty ()
std::vector< double > * get_counters ()
```

## 8.34.1 Detailed Description

Definition at line 44 of file geese-types.hpp.

## 8.34.2 Constructor & Destructor Documentation

## 8.34.2.1 PhyloCounterData() [1/2]

Definition at line 50 of file geese-types.hpp.

## 8.34.2.2 PhyloCounterData() [2/2]

```
PhyloCounterData::PhyloCounterData ( ) [inline]
```

Definition at line 55 of file geese-types.hpp.

## 8.34.3 Member Function Documentation

## 8.34.3.1 at()

Definition at line 57 of file geese-types.hpp.

## 8.34.3.2 begin()

```
std::vector< size_t >::iterator PhyloCounterData::begin ( ) [inline]
```

Definition at line 65 of file geese-types.hpp.

## 8.34.3.3 empty()

```
bool PhyloCounterData::empty ( ) [inline]
```

Definition at line 68 of file geese-types.hpp.

## 8.34.3.4 end()

```
std::vector< size_t >::iterator PhyloCounterData::end ( ) [inline]
```

Definition at line 66 of file geese-types.hpp.

## 8.34.3.5 get\_counters()

```
\verb|std::vector<| double >* PhyloCounterData::get_counters () [inline]| \\
```

Definition at line 69 of file geese-types.hpp.

#### 8.34.3.6 operator()()

Definition at line 58 of file geese-types.hpp.

## 8.34.3.7 operator[]()

Definition at line 59 of file geese-types.hpp.

#### 8.34.3.8 push back()

Definition at line 61 of file geese-types.hpp.

## 8.34.3.9 reserve()

Definition at line 60 of file geese-types.hpp.

## 8.34.3.10 shrink\_to\_fit()

```
void PhyloCounterData::shrink_to_fit ( ) [inline]
```

Definition at line 62 of file geese-types.hpp.

#### 8.34.3.11 size()

```
size_t PhyloCounterData::size ( ) [inline]
```

Definition at line 63 of file geese-types.hpp.

The documentation for this class was generated from the following file:

• include/barry/models/geese/geese-types.hpp

# 8.35 PhyloRuleDynData Class Reference

```
#include <geese-types.hpp>
```

#### **Public Member Functions**

- PhyloRuleDynData (const std::vector< double > \*counts\_, size\_t pos\_, size\_t lb\_, size\_t ub\_, size\_
   t duplication\_)
- const double operator() () const
- ∼PhyloRuleDynData ()

## **Public Attributes**

```
\bullet \ \ \mathsf{const} \ \mathsf{std} : \! \mathsf{vector} \! < \mathsf{double} > * \ \mathsf{counts}
```

```
    size t pos
```

- size\_t lb
- size\_t ub
- · size\_t duplication

## 8.35.1 Detailed Description

Definition at line 73 of file geese-types.hpp.

#### 8.35.2 Constructor & Destructor Documentation

## 8.35.2.1 PhyloRuleDynData()

Definition at line 81 of file geese-types.hpp.

## 8.35.2.2 ~PhyloRuleDynData()

```
PhyloRuleDynData::~PhyloRuleDynData ( ) [inline]
```

Definition at line 95 of file geese-types.hpp.

## 8.35.3 Member Function Documentation

## 8.35.3.1 operator()()

```
const double PhyloRuleDynData::operator() ( ) const [inline]
```

Definition at line 90 of file geese-types.hpp.

## 8.35.4 Member Data Documentation

#### 8.35.4.1 counts

```
const std::vector< double >* PhyloRuleDynData::counts
```

Definition at line 75 of file geese-types.hpp.

#### 8.35.4.2 duplication

```
size_t PhyloRuleDynData::duplication
```

Definition at line 79 of file geese-types.hpp.

## 8.35.4.3 lb

```
size_t PhyloRuleDynData::lb
```

Definition at line 77 of file geese-types.hpp.

#### 8.35.4.4 pos

```
size_t PhyloRuleDynData::pos
```

Definition at line 76 of file geese-types.hpp.

#### 8.35.4.5 ub

```
size_t PhyloRuleDynData::ub
```

Definition at line 78 of file geese-types.hpp.

The documentation for this class was generated from the following file:

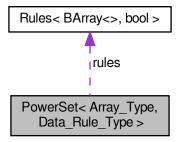
• include/barry/models/geese/geese-types.hpp

# 8.36 PowerSet< Array\_Type, Data\_Rule\_Type > Class Template Reference

Powerset of a binary array.

#include <powerset-bones.hpp>

Collaboration diagram for PowerSet < Array\_Type, Data\_Rule\_Type >:



#### **Public Member Functions**

- void init\_support ()
- void calc ()
- void reset (size\_t N\_, size\_t M\_)

## Construct and destroy a PowerSet object

- PowerSet ()
- PowerSet (size\_t N\_, size\_t M\_)
- PowerSet (const Array\_Type &array)
- ∼PowerSet ()

## Wrappers for the <tt>Rules</tt> member.

These will add rules to the model, which are shared by the support and the actual counter function.

- void add\_rule (Rule < Array\_Type, Data\_Rule\_Type > rule)
- void add\_rule (Rule\_fun\_type< Array\_Type, Data\_Rule\_Type > count\_fun\_, Data\_Rule\_Type data\_)

#### **Getter functions**

- const std::vector< Array\_Type > \* get\_data\_ptr () const
- std::vector< Array\_Type > get\_data () const
- std::vector< Array\_Type >::iterator begin ()
- std::vector< Array\_Type >::iterator end ()
- std::size\_t size () const noexcept
- const Array\_Type & operator[] (const size\_t &i) const

## **Public Attributes**

```
    Array_Type EmptyArray
```

- std::vector< Array\_Type > data
- Rules< Array\_Type, Data\_Rule\_Type > \* rules
- size t N
- size\_t M
- bool rules\_deleted = false
- std::vector< size\_t > coordinates\_free
- std::vector< size t > coordinates locked
- size t n free
- · size\_t n\_locked

## 8.36.1 Detailed Description

```
template<typename Array_Type = BArray<>>, typename Data_Rule_Type = bool> class PowerSet< Array_Type, Data_Rule_Type >
```

Powerset of a binary array.

**Template Parameters** 

Array_Type	
Data_Rule_Type	

Definition at line 11 of file powerset-bones.hpp.

## 8.36.2 Constructor & Destructor Documentation

#### 8.36.2.1 PowerSet() [1/3]

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
PowerSet< Array_Type, Data_Rule_Type >::PowerSet ( ) [inline]
```

Definition at line 36 of file powerset-bones.hpp.

#### 8.36.2.2 PowerSet() [2/3]

Definition at line 38 of file powerset-bones.hpp.

## 8.36.2.3 PowerSet() [3/3]

Definition at line 5 of file powerset-meat.hpp.

#### 8.36.2.4 ∼PowerSet()

```
template<typename Array_Type , typename Data_Rule_Type >
PowerSet< Array_Type, Data_Rule_Type >::~PowerSet [inline]
```

Definition at line 13 of file powerset-meat.hpp.

#### 8.36.3 Member Function Documentation

#### 8.36.3.1 add\_rule() [1/2]

Definition at line 180 of file powerset-meat.hpp.

#### 8.36.3.2 add\_rule() [2/2]

Definition at line 189 of file powerset-meat.hpp.

## 8.36.3.3 begin()

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< Array_Type >::iterator PowerSet< Array_Type, Data_Rule_Type >::begin ( ) [inline]
```

Definition at line 68 of file powerset-bones.hpp.

#### 8.36.3.4 calc()

```
template<typename Array_Type , typename Data_Rule_Type >
void PowerSet< Array_Type, Data_Rule_Type >::calc [inline]
```

Definition at line 151 of file powerset-meat.hpp.

#### 8.36.3.5 end()

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< Array_Type >::iterator PowerSet< Array_Type, Data_Rule_Type >::end ( ) [inline]
```

Definition at line 69 of file powerset-bones.hpp.

## 8.36.3.6 get\_data()

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< Array_Type > PowerSet< Array_Type, Data_Rule_Type >::get_data ( ) const [inline]
```

Definition at line 67 of file powerset-bones.hpp.

## 8.36.3.7 get\_data\_ptr()

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
const std::vector< Array_Type >* PowerSet< Array_Type, Data_Rule_Type >::get_data_ptr ()
const [inline]
```

Definition at line 66 of file powerset-bones.hpp.

## 8.36.3.8 init\_support()

```
template<typename Array_Type , typename Data_Rule_Type >
void PowerSet< Array_Type, Data_Rule_Type >::init_support [inline]
```

Definition at line 19 of file powerset-meat.hpp.

## 8.36.3.9 operator[]()

Definition at line 71 of file powerset-bones.hpp.

#### 8.36.3.10 reset()

Definition at line 167 of file powerset-meat.hpp.

#### 8.36.3.11 size()

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::size_t PowerSet< Array_Type, Data_Rule_Type >::size ( ) const [inline], [noexcept]
```

Definition at line 70 of file powerset-bones.hpp.

#### 8.36.4 Member Data Documentation

#### 8.36.4.1 coordinates\_free

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< size_t > PowerSet< Array_Type, Data_Rule_Type >::coordinates_free
```

Definition at line 26 of file powerset-bones.hpp.

#### 8.36.4.2 coordinates locked

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< size_t > PowerSet< Array_Type, Data_Rule_Type >::coordinates_locked
```

Definition at line 27 of file powerset-bones.hpp.

#### 8.36.4.3 data

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< Array_Type > PowerSet< Array_Type, Data_Rule_Type >::data
```

Definition at line 19 of file powerset-bones.hpp.

## 8.36.4.4 EmptyArray

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
Array_Type PowerSet< Array_Type, Data_Rule_Type >::EmptyArray
```

Definition at line 18 of file powerset-bones.hpp.

#### 8.36.4.5 M

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
size_t PowerSet< Array_Type, Data_Rule_Type >::M
```

Definition at line 22 of file powerset-bones.hpp.

#### 8.36.4.6 N

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
size_t PowerSet< Array_Type, Data_Rule_Type >::N
```

Definition at line 22 of file powerset-bones.hpp.

#### 8.36.4.7 n\_free

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
size_t PowerSet< Array_Type, Data_Rule_Type >::n_free
```

Definition at line 28 of file powerset-bones.hpp.

## 8.36.4.8 n\_locked

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
size_t PowerSet< Array_Type, Data_Rule_Type >::n_locked
```

Definition at line 29 of file powerset-bones.hpp.

#### 8.36.4.9 rules

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
Rules<Array_Type,Data_Rule_Type>* PowerSet< Array_Type, Data_Rule_Type >::rules
```

Definition at line 20 of file powerset-bones.hpp.

#### 8.36.4.10 rules\_deleted

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
bool PowerSet< Array_Type, Data_Rule_Type >::rules_deleted = false
```

Definition at line 23 of file powerset-bones.hpp.

The documentation for this class was generated from the following files:

- include/barry/powerset-bones.hpp
- include/barry/powerset-meat.hpp

## 8.37 Progress Class Reference

A simple progress bar.

```
#include progress.hpp>
```

## **Public Member Functions**

- Progress (int n\_, int width\_)
- ∼Progress ()
- void next ()
- void end ()

## 8.37.1 Detailed Description

A simple progress bar.

Definition at line 11 of file progress.hpp.

## 8.37.2 Constructor & Destructor Documentation

## 8.37.2.1 Progress()

Definition at line 30 of file progress.hpp.

## 8.37.2.2 ∼Progress()

```
Progress::~Progress ( ) [inline]
```

Definition at line 23 of file progress.hpp.

#### 8.37.3 Member Function Documentation

#### 8.37.3.1 end()

```
void Progress::end ( ) [inline]
```

Definition at line 52 of file progress.hpp.

## 8.37.3.2 next()

```
void Progress::next ( ) [inline]
```

Definition at line 41 of file progress.hpp.

The documentation for this class was generated from the following file:

• include/barry/progress.hpp

## 8.38 Rule < Array\_Type, Data\_Type > Class Template Reference

Rule for determining if a cell should be included in a sequence.

```
#include <rules-bones.hpp>
```

## **Public Member Functions**

- ∼Rule ()
- Data\_Type & D ()

Read/Write access to the data.

- bool operator() (const Array\_Type &a, size\_t i, size\_t j)
- std::string & get\_name ()
- std::string & get\_description ()
- std::string get\_name () const
- std::string get\_description () const

## Construct a new Rule object

Construct a new Rule object

#### **Parameters**

fun_	A function of type Rule_fun_type.
dat_	Data pointer to be passed to fun_
delete_←	When true, the Rule destructor will delete the pointer, if defined.
dat_	

- Rule ()
- Rule (Rule\_fun\_type< Array\_Type, Data\_Type > fun\_, Data\_Type dat\_, std::string name\_="", std::string desc ="")

## 8.38.1 Detailed Description

```
\label{template} $$ \ensuremath{\sf template}$$ < typename \ensuremath{\sf Array\_Type}$ = BArray<>, typename \ensuremath{\sf Data\_Type}$ = bool> class \ensuremath{\sf Rule}< Array\_Type, \ensuremath{\sf Data\_Type}>
```

Rule for determining if a cell should be included in a sequence.

Rules can be used together with Support and PowerSet to determine which cells should be included when enumerating all possible realizations of a binary array.

#### **Template Parameters**

Array_Type	An object of class BArray.
Data_Type	Any type.

Definition at line 20 of file rules-bones.hpp.

## 8.38.2 Constructor & Destructor Documentation

#### 8.38.2.1 Rule() [1/2]

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Rule< Array_Type, Data_Type >::Rule () [inline]
```

Definition at line 41 of file rules-bones.hpp.

## 8.38.2.2 Rule() [2/2]

Definition at line 42 of file rules-bones.hpp.

## 8.38.2.3 ∼Rule()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Rule< Array_Type, Data_Type >::~Rule ( ) [inline]
```

Definition at line 50 of file rules-bones.hpp.

#### 8.38.3 Member Function Documentation

## 8.38.3.1 D()

```
template<typename Array_Type , typename Data_Type >
Data_Type & Rule< Array_Type, Data_Type >::D [inline]
```

Read/Write access to the data.

Definition at line 37 of file rules-meat.hpp.

## 8.38.3.2 get\_description() [1/2]

```
template<typename Array_Type , typename Data_Type >
std::string & Rule< Array_Type, Data_Type >::get_description [inline]
```

Definition at line 54 of file rules-meat.hpp.

## 8.38.3.3 get\_description() [2/2]

```
template<typename Array_Type , typename Data_Type >
std::string Rule< Array_Type, Data_Type >::get_description [inline]
```

Definition at line 66 of file rules-meat.hpp.

## 8.38.3.4 get\_name() [1/2]

```
template<typename Array_Type , typename Data_Type >
std::string & Rule< Array_Type, Data_Type >::get_name [inline]
```

Definition at line 48 of file rules-meat.hpp.

#### 8.38.3.5 get\_name() [2/2]

```
template<typename Array_Type , typename Data_Type >
std::string Rule< Array_Type, Data_Type >::get_name [inline]
```

Definition at line 60 of file rules-meat.hpp.

#### 8.38.3.6 operator()()

Definition at line 43 of file rules-meat.hpp.

The documentation for this class was generated from the following files:

- include/barry/rules-bones.hpp
- include/barry/rules-meat.hpp

# 8.39 Rules < Array\_Type, Data\_Type > Class Template Reference

Vector of objects of class Rule.

```
#include <rules-bones.hpp>
```

#### **Public Member Functions**

- Rules ()
- Rules (const Rules < Array\_Type, Data\_Type > &rules\_)
- Rules< Array\_Type, Data\_Type > operator= (const Rules< Array\_Type, Data\_Type > &rules\_)
- ∼Rules ()
- size\_t size () const noexcept
- bool operator() (const Array\_Type &a, size\_t i, size\_t j)

Check whether a given cell is free or locked.

- void get\_seq (const Array\_Type &a, std::vector< size\_t > \*free, std::vector< size\_t > \*locked=nullptr)
   Computes the sequence of free and locked cells in an BArray.
- std::vector< std::string > get\_names () const
- std::vector< std::string > get\_descriptions () const
- std::vector< Rule< Array\_Type, Data\_Type > >::iterator begin ()
- std::vector< Rule< Array\_Type, Data\_Type > >::iterator end ()

#### Rule adding

#### **Parameters**

```
rule
```

- void add rule (Rule < Array Type, Data Type > rule)
- void add\_rule (Rule\_fun\_type< Array\_Type, Data\_Type > rule\_, Data\_Type data\_, std::string name\_="", std::string description\_="")

## 8.39.1 Detailed Description

```
template<typename Array_Type, typename Data_Type> class Rules< Array_Type, Data_Type >
```

Vector of objects of class Rule.

## **Template Parameters**

Array_Type	An object of class BArray
Data_Type	Any type.

Definition at line 71 of file rules-bones.hpp.

## 8.39.2 Constructor & Destructor Documentation

## 8.39.2.1 Rules() [1/2]

```
template<typename Array_Type , typename Data_Type >
Rules< Array_Type, Data_Type >::Rules ( ) [inline]
```

Definition at line 77 of file rules-bones.hpp.

## 8.39.2.2 Rules() [2/2]

Definition at line 5 of file rules-meat.hpp.

## 8.39.2.3 ∼Rules()

```
template<typename Array_Type , typename Data_Type >
Rules< Array_Type, Data_Type >::~Rules ( ) [inline]
```

Definition at line 82 of file rules-bones.hpp.

## 8.39.3 Member Function Documentation

#### 8.39.3.1 add rule() [1/2]

Definition at line 72 of file rules-meat.hpp.

#### 8.39.3.2 add rule() [2/2]

Definition at line 82 of file rules-meat.hpp.

## 8.39.3.3 begin()

```
template<typename Array_Type , typename Data_Type >
std::vector< Rule<Array_Type,Data_Type> >::iterator Rules< Array_Type, Data_Type >::begin (
) [inline]
```

Definition at line 134 of file rules-bones.hpp.

## 8.39.3.4 end()

```
template<typename Array_Type , typename Data_Type >
std::vector< Rule<Array_Type,Data_Type> >::iterator Rules< Array_Type, Data_Type >::end ( )
[inline]
```

Definition at line 137 of file rules-bones.hpp.

## 8.39.3.5 get\_descriptions()

```
template<typename Array_Type , typename Data_Type >
std::vector< std::string > Rules< Array_Type, Data_Type >::get_descriptions [inline]
```

Definition at line 180 of file rules-meat.hpp.

#### 8.39.3.6 get\_names()

```
template<typename Array_Type , typename Data_Type >
std::vector< std::string > Rules< Array_Type, Data_Type >::get_names [inline]
```

Definition at line 167 of file rules-meat.hpp.

## 8.39.3.7 get\_seq()

Computes the sequence of free and locked cells in an BArray.

#### **Parameters**

а	An object of class BArray.
free	Pointer to a vector of pairs (i, j) listing the free cells.
locked	(optional) Pointer to a vector of pairs (i, j) listing the locked cells.

#### Returns

Nothing.

Definition at line 117 of file rules-meat.hpp.

## 8.39.3.8 operator()()

Check whether a given cell is free or locked.

#### **Parameters**

а	A BArray object
i	row position
j	col position

#### Returns

true If the cell is locked false If the cell is free

Definition at line 101 of file rules-meat.hpp.

#### 8.39.3.9 operator=()

Definition at line 19 of file rules-meat.hpp.

## 8.39.3.10 size()

```
template<typename Array_Type , typename Data_Type >
size_t Rules< Array_Type, Data_Type >::size ( ) const [inline], [noexcept]
```

Definition at line 84 of file rules-bones.hpp.

The documentation for this class was generated from the following files:

- include/barry/rules-bones.hpp
- include/barry/rules-meat.hpp

# 8.40 StatsCounter< Array\_Type, Data\_Type > Class Template Reference

Count stats for a single Array.

```
#include <statscounter-bones.hpp>
```

#### **Public Member Functions**

```
    StatsCounter (const Array_Type *Array_)
```

Creator of a StatsCounter

StatsCounter (const StatsCounter < Array\_Type, Data\_Type > &counter)

Copy constructor.

• StatsCounter ()

Can be created without setting the array.

- ∼StatsCounter ()
- void reset\_array (const Array\_Type \*Array\_)

Changes the reference array for the counting.

- void add\_counter (Counter< Array\_Type, Data\_Type > f\_)
- void set\_counters (Counters < Array\_Type, Data\_Type > \*counters\_)
- void count\_init (size\_t i, size\_t j)

Counter functions This function recurses through the entries of Array and at each step of adding a new cell it uses the functions to list the statistics.

- void count\_current (size\_t i, size\_t j)
- std::vector< double > count\_all ()
- Counters< Array Type, Data Type > \* get counters ()
- std::vector< std::string > get\_names () const
- std::vector< std::string > get\_descriptions () const
- size t size () const

## 8.40.1 Detailed Description

```
\label{template} \begin{tabular}{ll} template < typename \ Array\_Type, \ typename \ Data\_Type > \\ class \ Stats Counter < Array\_Type, \ Data\_Type > \\ \end{tabular}
```

Count stats for a single Array.

Users can a list of functions that can be used with this. The baseline set of arguments is a pointer to a binary array and a dataset to add the counts to.

Definition at line 14 of file statscounter-bones.hpp.

#### 8.40.2 Constructor & Destructor Documentation

## 8.40.2.1 StatsCounter() [1/3]

Creator of a StatsCounter

#### **Parameters**

Array←	A const pointer to a BArray.

Definition at line 37 of file statscounter-bones.hpp.

## 8.40.2.2 StatsCounter() [2/3]

Copy constructor.

#### **Parameters**

counter

## 8.40.2.3 StatsCounter() [3/3]

```
template<typename Array_Type , typename Data_Type >
StatsCounter< Array_Type, Data_Type >::StatsCounter ( ) [inline]
```

Can be created without setting the array.

Definition at line 59 of file statscounter-bones.hpp.

## 8.40.2.4 $\sim$ StatsCounter()

```
template<typename Array_Type , typename Data_Type >
StatsCounter< Array_Type, Data_Type >::~StatsCounter ( )
```

## 8.40.3 Member Function Documentation

## 8.40.3.1 add\_counter()

## 8.40.3.2 count\_all()

```
template<typename Array_Type , typename Data_Type >
std::vector< double > StatsCounter< Array_Type, Data_Type >::count_all [inline]
```

Definition at line 99 of file statscounter-meat.hpp.

## 8.40.3.3 count\_current()

## 8.40.3.4 count\_init()

Counter functions This function recurses through the entries of Array and at each step of adding a new cell it uses the functions to list the statistics.

## 8.40.3.5 get\_counters()

```
template<typename Array_Type , typename Data_Type >
Counters<Array_Type,Data_Type>* StatsCounter< Array_Type, Data_Type >::get_counters ( )
```

## 8.40.3.6 get\_descriptions()

```
template<typename Array_Type , typename Data_Type >
std::vector< std::string > StatsCounter< Array_Type, Data_Type >::get_descriptions ( ) const
```

#### 8.40.3.7 get\_names()

```
template<typename Array_Type , typename Data_Type >
std::vector< std::string > StatsCounter< Array_Type, Data_Type >::get_names ( ) const
```

## 8.40.3.8 reset\_array()

Changes the reference array for the counting.

#### **Parameters**

Array⇔	A pointer to an array of class Array_Type.

#### 8.40.3.9 set counters()

#### 8.40.3.10 size()

```
template<typename Array_Type , typename Data_Type >
size_t StatsCounter< Array_Type, Data_Type >::size ( ) const [inline]
```

Definition at line 86 of file statscounter-bones.hpp.

The documentation for this class was generated from the following files:

- include/barry/statscounter-bones.hpp
- include/barry/statscounter-meat.hpp

# 8.41 Support < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type > Class Template Reference

Compute the support of sufficient statistics.

```
#include <support-bones.hpp>
```

## **Public Member Functions**

Support (const Array\_Type &Array\_)

Constructor passing a reference Array.

• Support (size\_t N\_, size\_t M\_)

Constructor specifying the dimensions of the array (empty).

- Support ()
- ∼Support ()
- void init\_support (std::vector< Array\_Type > \*array\_bank=nullptr, std::vector< double > \*stats\_\cup bank=nullptr)
- void calc (std::vector< Array\_Type > \*array\_bank=nullptr, std::vector< double > \*stats\_bank=nullptr, size
   \_t max\_num\_elements\_=0u)

Computes the entire support.

- const std::vector< double > & get\_counts () const
- std::vector< double > \* get\_current\_stats ()

List current statistics.

- void print () const
- const FreqTable< double > & get\_data () const
- Counters < Array\_Type, Data\_Counter\_Type > \* get\_counters ()

Vector of couter functions.

Rules< Array\_Type, Data\_Rule\_Type > \* get\_rules ()

Vector of static rules (cells to iterate).

Rules< Array\_Type, Data\_Rule\_Dyn\_Type > \* get\_rules\_dyn ()

Vector of dynamic rules (to include/exclude a realizaton).

## Resets the support calculator

If needed, the counters of a support object can be reused.

#### **Parameters**

Array←	New array over which the support will be computed.

- void reset array ()
- void reset\_array (const Array\_Type &Array\_)

#### Manage counters

#### **Parameters**

f_	A counter to be added.
counters←	A vector of counters to be added.

- void add\_counter (Counter< Array\_Type, Data\_Counter\_Type > f\_)
- void set\_counters (Counters < Array\_Type, Data\_Counter\_Type > \*counters\_)

#### Manage rules

#### **Parameters**

f_	A rule to be added.
counters←	A vector of rules to be added.
_	

- void add\_rule (Rule < Array\_Type, Data\_Rule\_Type > \*f\_)
- void add\_rule (Rule < Array\_Type, Data\_Rule\_Type > f\_)
- void set\_rules (Rules< Array\_Type, Data\_Rule\_Type > \*rules\_)
- void add\_rule\_dyn (Rule < Array\_Type, Data\_Rule\_Dyn\_Type > \*f\_)
- void add\_rule\_dyn (Rule< Array\_Type, Data\_Rule\_Dyn\_Type > f\_)
- void set\_rules\_dyn (Rules < Array\_Type, Data\_Rule\_Dyn\_Type > \*rules\_)
- bool eval\_rules\_dyn (const std::vector< double > &counts, const size\_t &i, const size\_t &j)

#### **Public Attributes**

- size t N
- size\_t M
- bool delete\_counters = true
- bool delete rules = true
- bool delete rules dyn = true
- size t max num elements = BARRY MAX NUM ELEMENTS
- std::vector< double > current\_stats
- std::vector< size\_t > coordinates\_free
- std::vector< size t > coordinates locked
- · size t coordiantes n free
- size\_t coordiantes\_n\_locked
- std::vector< double > change stats
- std::vector< size t > hashes
- std::vector< bool > hashes\_initialized
- size\_t n\_counters

## 8.41.1 Detailed Description

template < typename Array\_Type = BArray < bool, bool >, typename Data\_Counter\_Type = bool, typename Data\_Rule\_Type = bool, typename Data\_Rule\_Dyn\_Type = bool > class Support < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Dyn\_Type >

Compute the support of sufficient statistics.

Given an array and a set of counters, this object iterates throughout the support set of the Array while at the same time computing the support of the sufficient statitics.

The members rule and rule\_dyn allow constraining the support. The first will establish which cells of the array will be used to iterate, for example, in the case of social networks, self-loops are not allowed, so the entire diagonal would be fixed to zero, reducing the size of the support.

In the case of rule\_dyn, the function will stablish dynamically whether the current state will be included in the counts or not. For example, this set of rules can be used to constrain the support to networks that have a prescribed degree sequence.

Definition at line 42 of file support-bones.hpp.

## 8.41.2 Constructor & Destructor Documentation

## 8.41.2.1 Support() [1/3]

Constructor passing a reference Array.

Definition at line 89 of file support-bones.hpp.

## 8.41.2.2 Support() [2/3]

Constructor specifying the dimensions of the array (empty).

Definition at line 98 of file support-bones.hpp.

## 8.41.2.3 Support() [3/3]

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename
Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::Support ( )
[inline]
```

Definition at line 105 of file support-bones.hpp.

#### 8.41.2.4 ∼Support()

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename
Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::~Support ()
[inline]
```

Definition at line 112 of file support-bones.hpp.

#### 8.41.3 Member Function Documentation

## 8.41.3.1 add\_counter()

Definition at line 417 of file support-meat.hpp.

## 8.41.3.2 add\_rule() [1/2]

Definition at line 444 of file support-meat.hpp.

## 8.41.3.3 add\_rule() [2/2]

Definition at line 454 of file support-meat.hpp.

#### 8.41.3.4 add\_rule\_dyn() [1/2]

Definition at line 479 of file support-meat.hpp.

## 8.41.3.5 add\_rule\_dyn() [2/2]

Definition at line 489 of file support-meat.hpp.

## 8.41.3.6 calc()

Computes the entire support.

Not to be used by the user. Sets the starting point in the array (column-major).

#### **Parameters**

array_bank	If specified, the counter will add to the vector each possible state of the array, as it counts.
stats_bank	If specified, the counter will add to the vector each possible set of statistics, as it counts.

Definition at line 383 of file support-meat.hpp.

#### 8.41.3.7 eval\_rules\_dyn()

Definition at line 514 of file support-meat.hpp.

## 8.41.3.8 get\_counters()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >

Counters< Array_Type, Data_Counter_Type > * Support< Array_Type, Data_Counter_Type, Data_←

Rule_Type, Data_Rule_Dyn_Type >::get_counters [inline]
```

Vector of couter functions.

Definition at line 593 of file support-meat.hpp.

#### 8.41.3.9 get\_counts()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type > const std::vector< double > & Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_\leftarrow Rule_Dyn_Type >::get_counts [inline]
```

Definition at line 557 of file support-meat.hpp.

## 8.41.3.10 get\_current\_stats()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< double > * Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_\Lorentype, Dyn_Type >::get_current_stats [inline]
```

List current statistics.

Definition at line 571 of file support-meat.hpp.

#### 8.41.3.11 get\_data()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
const FreqTable< double > & Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_←
Rule_Dyn_Type >::get_data [inline]
```

Definition at line 588 of file support-meat.hpp.

## 8.41.3.12 get\_rules()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
Rules< Array_Type, Data_Rule_Type > * Support< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_rules [inline]
```

Vector of static rules (cells to iterate).

Definition at line 598 of file support-meat.hpp.

## 8.41.3.13 get\_rules\_dyn()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >

Rules< Array_Type, Data_Rule_Dyn_Type > * Support< Array_Type, Data_Counter_Type, Data_Rule

_Type, Data_Rule_Dyn_Type >::get_rules_dyn [inline]
```

Vector of dynamic rules (to include/exclude a realizaton).

Definition at line 603 of file support-meat.hpp.

## 8.41.3.14 init\_support()

Definition at line 5 of file support-meat.hpp.

#### 8.41.3.15 print()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::print
[inline]
```

Definition at line 576 of file support-meat.hpp.

## 8.41.3.16 reset\_array() [1/2]

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::reset_array
[inline]
```

Definition at line 111 of file support-meat.hpp.

## 8.41.3.17 reset\_array() [2/2]

Definition at line 118 of file support-meat.hpp.

## 8.41.3.18 set\_counters()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >

void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::set_

counters (

Counters< Array_Type, Data_Counter_Type > * counters_) [inline]
```

Definition at line 427 of file support-meat.hpp.

## 8.41.3.19 set\_rules()

Definition at line 464 of file support-meat.hpp.

## 8.41.3.20 set\_rules\_dyn()

Definition at line 499 of file support-meat.hpp.

#### 8.41.4 Member Data Documentation

## 8.41.4.1 change\_stats

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< double > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn \( \to \) _Type >::change_stats
```

Definition at line 82 of file support-bones.hpp.

## 8.41.4.2 coordiantes\_n\_free

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::coordiantes 
n free
```

Definition at line 80 of file support-bones.hpp.

#### 8.41.4.3 coordiantes\_n\_locked

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::coordiantes \cdot _n_locked
```

Definition at line 81 of file support-bones.hpp.

#### 8.41.4.4 coordinates free

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< size_t > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn 
_Type >::coordinates_free
```

Definition at line 78 of file support-bones.hpp.

#### 8.41.4.5 coordinates\_locked

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< size_t > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn 
_Type >::coordinates_locked
```

Definition at line 79 of file support-bones.hpp.

## 8.41.4.6 current\_stats

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< double > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn← _Type >::current_stats
```

Definition at line 77 of file support-bones.hpp.

## 8.41.4.7 delete\_counters

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_← counters = true
```

Definition at line 71 of file support-bones.hpp.

#### 8.41.4.8 delete\_rules

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_← rules = true
```

Definition at line 72 of file support-bones.hpp.

#### 8.41.4.9 delete rules dyn

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_← rules_dyn = true
```

Definition at line 73 of file support-bones.hpp.

#### 8.41.4.10 hashes

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< size_t > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn← _Type >::hashes
```

Definition at line 83 of file support-bones.hpp.

## 8.41.4.11 hashes\_initialized

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< bool > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_↔ Type >::hashes_initialized
```

Definition at line 84 of file support-bones.hpp.

#### 8.41.4.12 M

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename
Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::M
```

Definition at line 70 of file support-bones.hpp.

#### 8.41.4.13 max\_num\_elements

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename
Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::max_num
_elements = BARRY_MAX_NUM_ELEMENTS
```

Definition at line 74 of file support-bones.hpp.

#### 8.41.4.14 N

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename
Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::N
```

Definition at line 70 of file support-bones.hpp.

#### 8.41.4.15 n counters

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::n_
counters
```

Definition at line 85 of file support-bones.hpp.

The documentation for this class was generated from the following files:

- include/barry/support-bones.hpp
- include/barry/support-meat.hpp

## 8.42 vecHasher < T > Struct Template Reference

#include <typedefs.hpp>

## **Public Member Functions**

• std::size\_t operator() (std::vector< T > const &dat) const noexcept

## 8.42.1 Detailed Description

```
\label{eq:typename} \begin{array}{l} \text{template}\!<\!\text{typename T}\!>\\ \text{struct vecHasher}\!<\!\text{T}\!> \end{array}
```

Definition at line 105 of file typedefs.hpp.

## 8.42.2 Member Function Documentation

## 8.42.2.1 operator()()

Definition at line 108 of file typedefs.hpp.

The documentation for this struct was generated from the following file:

• include/barry/typedefs.hpp

# **Chapter 9**

# **File Documentation**

# 9.1 include/barry/barray-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



## **Classes**

class BArray < Cell\_Type, Data\_Type >
 Baseline class for binary arrays.

# 9.2 include/barry/barray-iterator.hpp File Reference

## Classes

class ConstBArrayRowIter< Cell\_Type, Data\_Type >

252 File Documentation

## 9.3 include/barry/barray-meat-operators.hpp File Reference

This graph shows which files directly or indirectly include this file:



## **Macros**

- #define BARRAY\_TYPE() BArray<Cell\_Type, Data\_Type>
- #define BARRAY\_TEMPLATE\_ARGS() < typename Cell\_Type, typename Data\_Type>
- #define BARRAY\_TEMPLATE(a, b) template BARRAY\_TEMPLATE\_ARGS() inline a BARRAY\_TYPE()::b
- #define ROW(a) this->el\_ij[a]
- #define COL(a) this->el\_ji[a]

#### **Functions**

- template BARRAY TEMPLATE ARGS () inline void checkdim (const BARRAY TYPE() &lhs
- template const BARRAY TYPE () &rhs)
- BARRAY\_TEMPLATE (BARRAY\_TYPE()&, operator+=)(const BArray< Cell\_Type
- for (size\_t i=0u;i< nrow();++i) for(size\_t j=0u
- j< ncol();++j) this-> operator() (i, j)+
- BARRAY\_TEMPLATE (BARRAY\_TYPE()&, operator+=)(const Cell\_Type &rhs)
- BARRAY\_TEMPLATE (BARRAY\_TYPE()&, operator-=)(const BArray< Cell\_Type
- BARRAY\_TEMPLATE (BARRAY\_TYPE()&, operator-=)(const Cell\_Type &rhs)
- BARRAY\_TEMPLATE (BARRAY\_TYPE()&, operator\*=)(const Cell\_Type &rhs)
- BARRAY\_TEMPLATE (BARRAY\_TYPE()&, operator/=)(const Cell\_Type &rhs)

## **Variables**

- Data\_Type & rhs
- return \* this

#### 9.3.1 Macro Definition Documentation

## 9.3.1.1 BARRAY\_TEMPLATE

Definition at line 11 of file barray-meat-operators.hpp.

## 9.3.1.2 BARRAY\_TEMPLATE\_ARGS

```
#define BARRAY_TEMPLATE_ARGS( ) <typename Cell_Type, typename Data_Type>
```

Definition at line 9 of file barray-meat-operators.hpp.

## 9.3.1.3 BARRAY\_TYPE

```
#define BARRAY_TYPE( ) BArray<Cell_Type, Data_Type>
```

Definition at line 7 of file barray-meat-operators.hpp.

## 9.3.1.4 COL

Definition at line 15 of file barray-meat-operators.hpp.

#### 9.3.1.5 ROW

Definition at line 14 of file barray-meat-operators.hpp.

## 9.3.2 Function Documentation

## 9.3.2.1 BARRAY\_TEMPLATE() [1/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator* ) const &
```

Definition at line 88 of file barray-meat-operators.hpp.

## 9.3.2.2 BARRAY\_TEMPLATE() [2/6]

## 9.3.2.3 BARRAY\_TEMPLATE() [3/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator+ ) const &
```

Definition at line 46 of file barray-meat-operators.hpp.

## 9.3.2.4 BARRAY\_TEMPLATE() [4/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator- ) const
```

## 9.3.2.5 BARRAY\_TEMPLATE() [5/6]

Definition at line 75 of file barray-meat-operators.hpp.

## **9.3.2.6 BARRAY\_TEMPLATE()** [6/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator/ ) const &
```

Definition at line 105 of file barray-meat-operators.hpp.

## 9.3.2.7 BARRAY\_TEMPLATE\_ARGS()

```
template BARRAY_TEMPLATE_ARGS ( ) const \&
```

## 9.3.2.8 BARRAY\_TYPE()

```
template const BARRAY_TYPE ( ) &
```

Definition at line 20 of file barray-meat-operators.hpp.

#### 9.3.2.9 for()

```
for ( ) [pure virtual]
```

Definition at line 66 of file barray-meat-operators.hpp.

## 9.3.2.10 operator()()

## 9.3.3 Variable Documentation

#### 9.3.3.1 rhs

```
Data_Type & rhs
Initial value:
{
    checkdim_(*this, rhs)
```

Definition at line 33 of file barray-meat-operators.hpp.

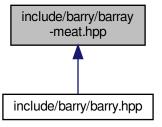
#### 9.3.3.2 this

```
return * this
```

Definition at line 43 of file barray-meat-operators.hpp.

# 9.4 include/barry/barray-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



## **Macros**

- #define ROW(a) this->el\_ij[a]
- #define COL(a) this->el\_ji[a]

### 9.4.1 Macro Definition Documentation

#### 9.4.1.1 COL

Definition at line 14 of file barray-meat.hpp.

#### 9.4.1.2 ROW

Definition at line 13 of file barray-meat.hpp.

# 9.5 include/barry/barraycell-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



## **Classes**

- class BArrayCell
   Cell\_Type, Data\_Type
- class BArrayCell\_const< Cell\_Type, Data\_Type >

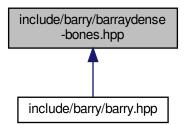
# 9.6 include/barry/barraycell-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



# 9.7 include/barry/barraydense-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



## Classes

class BArrayDense < Cell\_Type, Data\_Type >
 Baseline class for binary arrays.

## 9.8 include/barry/barraydense-meat-operators.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Macros**

- #define BDENSE\_TYPE() BArrayDense<Cell\_Type, Data\_Type>
- $\bullet \ \ \text{\#define BDENSE\_TEMPLATE\_ARGS()} < \text{typename Cell\_Type, typename Data\_Type} >$
- #define BDENSE\_TEMPLATE(a, b) template BDENSE\_TEMPLATE\_ARGS() inline a BDENSE\_TYPE()::b
- #define ROW(a) this->el ij[a]
- #define COL(a) this->el\_ji[a]
- #define POS(a, b) (b)\*N + (a)
- #define POS\_N(a, b, c) (b)\*(c) + (a)

#### **Functions**

- template BDENSE TEMPLATE ARGS () inline void checkdim (const BDENSE TYPE() &lhs
- template const BDENSE TYPE () &rhs)
- BDENSE\_TEMPLATE (BDENSE\_TYPE()&, operator+=)(const BDENSE\_TYPE() &rhs)
- BDENSE\_TEMPLATE (BDENSE\_TYPE()&, operator-=)(const BDENSE\_TYPE() &rhs)
- BDENSE\_TEMPLATE (BDENSE\_TYPE()&, operator\*=)(const Cell\_Type &rhs)
- BDENSE\_TEMPLATE (BDENSE\_TYPE()&, operator/=)(const Cell\_Type &rhs)

#### 9.8.1 Macro Definition Documentation

#### 9.8.1.1 BDENSE\_TEMPLATE

Definition at line 11 of file barraydense-meat-operators.hpp.

## 9.8.1.2 BDENSE\_TEMPLATE\_ARGS

```
#define BDENSE_TEMPLATE_ARGS( ) <typename Cell_Type, typename Data_Type>
```

Definition at line 9 of file barraydense-meat-operators.hpp.

## 9.8.1.3 BDENSE\_TYPE

```
#define BDENSE_TYPE( ) BArrayDense<Cell_Type, Data_Type>
```

Definition at line 7 of file barraydense-meat-operators.hpp.

#### 9.8.1.4 COL

Definition at line 15 of file barraydense-meat-operators.hpp.

#### 9.8.1.5 POS

```
#define POS(  a, \\ b ) \ (b)*N + (a)
```

Definition at line 16 of file barraydense-meat-operators.hpp.

## 9.8.1.6 POS\_N

Definition at line 17 of file barraydense-meat-operators.hpp.

#### 9.8.1.7 ROW

Definition at line 14 of file barraydense-meat-operators.hpp.

#### 9.8.2 Function Documentation

## 9.8.2.1 BDENSE\_TEMPLATE() [1/4]

Definition at line 90 of file barraydense-meat-operators.hpp.

## 9.8.2.2 BDENSE\_TEMPLATE() [2/4]

Definition at line 34 of file barraydense-meat-operators.hpp.

#### 9.8.2.3 BDENSE\_TEMPLATE() [3/4]

Definition at line 61 of file barraydense-meat-operators.hpp.

## 9.8.2.4 BDENSE\_TEMPLATE() [4/4]

Definition at line 101 of file barraydense-meat-operators.hpp.

## 9.8.2.5 BDENSE\_TEMPLATE\_ARGS()

```
template BDENSE_TEMPLATE_ARGS ( ) const &
```

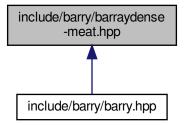
## 9.8.2.6 BDENSE\_TYPE()

```
template const BDENSE_TYPE ( ) &
```

Definition at line 22 of file barraydense-meat-operators.hpp.

## 9.9 include/barry/barraydense-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



### **Macros**

```
• #define ROW(a) this->el_ij[a]
```

- #define COL(a) this->el\_ji[a]
- #define POS(a, b) (b)\*N + (a)
- #define POS\_N(a, b, c) (b)\*(c) + (a)
- #define ZERO\_CELL static\_cast<Cell\_Type>(0.0)

#### 9.9.1 Macro Definition Documentation

#### 9.9.1.1 COL

Definition at line 24 of file barraydense-meat.hpp.

#### 9.9.1.2 POS

```
#define POS(  a, \\ b ) \ (b)*N + (a)
```

Definition at line 25 of file barraydense-meat.hpp.

## 9.9.1.3 POS\_N

Definition at line 26 of file barraydense-meat.hpp.

#### 9.9.1.4 ROW

Definition at line 23 of file barraydense-meat.hpp.

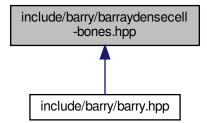
## 9.9.1.5 ZERO\_CELL

```
#define ZERO_CELL static_cast<Cell_Type>(0.0)
```

Definition at line 31 of file barraydense-meat.hpp.

# 9.10 include/barry/barraydensecell-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Classes**

- class BArrayDenseCell< Cell\_Type, Data\_Type >

#### **Macros**

```
• #define POS(a, b) (a) + (b) * N
```

## 9.10.1 Macro Definition Documentation

#### 9.10.1.1 POS

```
#define POS(  a, \\ b \ ) \ (a) \ + \ (b) \ * \ \mbox{N}
```

Definition at line 6 of file barraydensecell-bones.hpp.

# 9.11 include/barry/barraydensecell-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Macros**

```
• #define POS(a, b) (a) + (b) * dat->N
```

## 9.11.1 Macro Definition Documentation

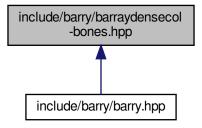
#### 9.11.1.1 POS

```
#define POS(  a, \\ b ) (a) + (b) * dat->N
```

Definition at line 6 of file barraydensecell-meat.hpp.

## 9.12 include/barry/barraydensecol-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Classes**

- class BArrayDenseCol< Cell\_Type, Data\_Type >
- class BArrayDenseCol\_const< Cell\_Type, Data\_Type >

## **Macros**

- #define POS(a, b) (b)\*N + (a)
- #define POS\_N(a, b, c) (b)\*(c) + (a)
- #define ZERO\_CELL static\_cast<Cell\_Type>(0.0)

#### 9.12.1 Macro Definition Documentation

### 9.12.1.1 POS

```
#define POS(  a, \\ b ) (b)*N + (a)
```

Definition at line 4 of file barraydensecol-bones.hpp.

## 9.12.1.2 POS\_N

Definition at line 5 of file barraydensecol-bones.hpp.

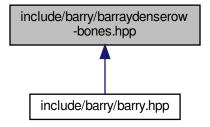
## 9.12.1.3 ZERO\_CELL

```
#define ZERO_CELL static_cast<Cell_Type>(0.0)
```

Definition at line 6 of file barraydensecol-bones.hpp.

# 9.13 include/barry/barraydenserow-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



## **Classes**

- class BArrayDenseRow
   Cell\_Type, Data\_Type >
- class BArrayDenseRow\_const< Cell\_Type, Data\_Type >

## **Macros**

- #define POS(a, b) (b) \* N + (a)
- #define POS\_N(a, b, c) (b)\*(c) + (a)
- #define ZERO\_CELL static\_cast< Cell\_Type >(0.0)

#### 9.13.1 Macro Definition Documentation

#### 9.13.1.1 POS

```
#define POS(  a, \\ b ) \ (b) \ * \ N \ + \ (a)
```

Definition at line 4 of file barraydenserow-bones.hpp.

#### 9.13.1.2 POS N

Definition at line 5 of file barraydenserow-bones.hpp.

#### 9.13.1.3 ZERO\_CELL

```
#define ZERO_CELL static_cast< Cell_Type >(0.0)
```

Definition at line 6 of file barraydenserow-bones.hpp.

## 9.14 include/barry/barrayrow-bones.hpp File Reference

#### **Classes**

- class BArrayRow
   Cell\_Type, Data\_Type >
- class BArrayRow const< Cell Type, Data Type >

## 9.15 include/barry/barrayrow-meat.hpp File Reference

## **Macros**

- #define BROW\_TYPE() BArrayRow<Cell\_Type, Data\_Type>
- #define BROW\_TEMPLATE\_ARGS() < typename Cell\_Type, typename Data\_Type >
- #define BROW\_TEMPLATE(a, b) template BROW\_TEMPLATE\_ARGS() inline a BROW\_TYPE()::b

#### **Functions**

- BROW\_TEMPLATE (void, operator=)(const BROW\_TYPE() &val)
- BROW\_TEMPLATE (void, operator+=)(const BROW\_TYPE() &val)
- BROW\_TEMPLATE (void, operator-=)(const BROW\_TYPE() &val)
- BROW\_TEMPLATE (void, operator\*=)(const BROW\_TYPE() &val)
- BROW\_TEMPLATE (void, operator/=)(const BROW\_TYPE() &val)

## 9.15.1 Macro Definition Documentation

#### 9.15.1.1 BROW\_TEMPLATE

Definition at line 8 of file barrayrow-meat.hpp.

#### 9.15.1.2 BROW\_TEMPLATE\_ARGS

```
#define BROW_TEMPLATE_ARGS( ) <typename Cell_Type, typename Data_Type>
```

Definition at line 6 of file barrayrow-meat.hpp.

## 9.15.1.3 BROW\_TYPE

```
#define BROW_TYPE( ) BArrayRow<Cell_Type, Data_Type>
```

Definition at line 4 of file barrayrow-meat.hpp.

## 9.15.2 Function Documentation

### 9.15.2.1 BROW\_TEMPLATE() [1/5]

Definition at line 45 of file barrayrow-meat.hpp.

## 9.15.2.2 BROW\_TEMPLATE() [2/5]

Definition at line 25 of file barrayrow-meat.hpp.

## 9.15.2.3 BROW\_TEMPLATE() [3/5]

Definition at line 34 of file barrayrow-meat.hpp.

#### 9.15.2.4 BROW\_TEMPLATE() [4/5]

Definition at line 55 of file barrayrow-meat.hpp.

## 9.15.2.5 BROW\_TEMPLATE() [5/5]

Definition at line 11 of file barrayrow-meat.hpp.

# 9.16 include/barry/barrayvector-bones.hpp File Reference

#### **Classes**

```
    class BArrayVector< Cell_Type, Data_Type >
    Row or column of a BArray
```

class BArrayVector\_const< Cell\_Type, Data\_Type >

## 9.17 include/barry/barrayvector-meat.hpp File Reference

# 9.18 include/barry/barry-configuration.hpp File Reference

This graph shows which files directly or indirectly include this file:



## **Configuration MACROS**

These are mostly related to performance. The definitions follow:

- BARRY\_USE\_UNORDERED\_MAP If specified, then barry is compiled using std::unordered\_map. Otherwise it will use std::map for the arrays.
- BARRY\_USE\_SAFE\_EXP When specified, it will multiply all likelihoods in Model by (1/-100)/(1/-100) so that numerical overflows are avoided.
- BARRY\_USE\_ISFINITE When specified, it will introduce a macro that checks whether the likelihood is finite or not.
- $printf\_barry$  If not specified, will be defined as printf.
- ${\tt BARRY\_DEBUG\_LEVEL},$  when defined, will make things verbose.
- #define BARRY\_SAFE\_EXP -100.0
- #define BARRY ISFINITE(a)
- #define BARRY\_CHECK\_SUPPORT(x, maxs)
- #define printf\_barry printf
- #define BARRY\_MAX\_NUM\_ELEMENTS static\_cast< size\_t >(std::numeric\_limits< size\_t >::max() /2u)
- template<typename Ta , typename Tb > using Map = std::map< Ta, Tb >

#### 9.18.1 Macro Definition Documentation

#### 9.18.1.1 BARRY CHECK SUPPORT

```
#define BARRY_CHECK_SUPPORT(
          x,
          maxs )
```

Definition at line 47 of file barry-configuration.hpp.

#### 9.18.1.2 BARRY\_ISFINITE

Definition at line 40 of file barry-configuration.hpp.

## 9.18.1.3 BARRY\_MAX\_NUM\_ELEMENTS

```
#define BARRY_MAX_NUM_ELEMENTS static_cast< size_t >(std::numeric_limits< size_t >::max()
/2u)
```

Definition at line 55 of file barry-configuration.hpp.

#### 9.18.1.4 BARRY\_SAFE\_EXP

```
#define BARRY_SAFE_EXP -100.0
```

Definition at line 33 of file barry-configuration.hpp.

## 9.18.1.5 printf\_barry

```
#define printf_barry printf
```

Definition at line 51 of file barry-configuration.hpp.

## 9.18.2 Typedef Documentation

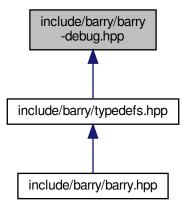
#### 9.18.2.1 Map

```
template<typename Ta , typename Tb >
using Map = std::map<Ta,Tb>
```

Definition at line 27 of file barry-configuration.hpp.

## 9.19 include/barry/barry-debug.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Macros**

• #define BARRY\_DEBUG\_LEVEL 0

## 9.19.1 Macro Definition Documentation

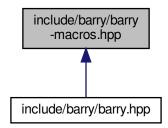
## 9.19.1.1 BARRY\_DEBUG\_LEVEL

#define BARRY\_DEBUG\_LEVEL 0

Definition at line 5 of file barry-debug.hpp.

## 9.20 include/barry/barry-macros.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Macros**

- #define BARRY\_ZERO Cell
   Cell\_Type>(0.0)
- #define BARRY\_ZERO\_DENSE static\_cast<Cell\_Type>(0.0)
- #define BARRY\_ONE Cell<Cell\_Type>(1.0)
- #define BARRY\_ONE\_DENSE static\_cast<Cell\_Type>(1.0)
- #define BARRY\_UNUSED(expr) do { (void)(expr); } while (0);
- #define BARRY\_NCORES\_ARG(default) size\_t ncores default

#### 9.20.1 Macro Definition Documentation

#### 9.20.1.1 BARRY\_NCORES\_ARG

```
\begin{tabular}{ll} \# define $BARRY_NCORES\_ARG($$ $default ) $size\_t $ncores $default$ \\ \end{tabular}
```

Definition at line 15 of file barry-macros.hpp.

## 9.20.1.2 BARRY\_ONE

```
#define BARRY_ONE CellCell_Type>(1.0)
```

Definition at line 7 of file barry-macros.hpp.

## 9.20.1.3 BARRY\_ONE\_DENSE

```
#define BARRY_ONE_DENSE static_cast<Cell_Type>(1.0)
```

Definition at line 8 of file barry-macros.hpp.

#### 9.20.1.4 BARRY\_UNUSED

Definition at line 10 of file barry-macros.hpp.

#### 9.20.1.5 BARRY\_ZERO

```
#define BARRY_ZERO Cell<Cell_Type>(0.0)
```

Definition at line 4 of file barry-macros.hpp.

#### 9.20.1.6 BARRY\_ZERO\_DENSE

```
#define BARRY_ZERO_DENSE static_cast<Cell_Type>(0.0)
```

Definition at line 5 of file barry-macros.hpp.

## 9.21 include/barry/barry.hpp File Reference

```
#include <iostream>
#include <cstdarg>
#include <vector>
#include <unordered_map>
#include <functional>
#include <stdexcept>
#include <cmath>
#include <map>
#include <algorithm>
#include <utility>
#include <random>
#include <climits>
#include <cfloat>
#include <string>
#include <cstdint>
#include <memory>
#include <regex>
```

```
#include <iterator>
#include "typedefs.hpp"
#include "barry-macros.hpp"
#include "freqtable.hpp"
#include "cell-bones.hpp"
#include "cell-meat.hpp"
#include "barray-bones.hpp"
#include "barraycell-bones.hpp"
#include "barray-meat.hpp"
#include "barraycell-meat.hpp"
#include "barray-meat-operators.hpp"
#include "barraydense-bones.hpp"
#include "barraydensecell-bones.hpp"
#include "barraydenserow-bones.hpp"
#include "barraydensecol-bones.hpp"
#include "barraydense-meat.hpp"
#include "barraydensecell-meat.hpp"
#include "barraydense-meat-operators.hpp"
#include "counters-bones.hpp"
#include "counters-meat.hpp"
#include "statscounter-bones.hpp"
#include "statscounter-meat.hpp"
#include "support-bones.hpp"
#include "support-meat.hpp"
#include "powerset-bones.hpp"
#include "powerset-meat.hpp"
#include "model-bones.hpp"
#include "model-meat.hpp"
#include "rules-bones.hpp"
#include "rules-meat.hpp"
#include "counters/network.hpp"
Include dependency graph for barry.hpp:
```

#### **Namespaces**

barry

barry: Your go-to motif accountant

barry::counters

Tree class and Treelterator class.

· barry::counters::network

#### **Macros**

- #define BARRY HPP
- #define BARRY VERSION MAYOR 0
- #define BARRY\_VERSION\_MINOR 1
- #define BARRY\_VERSION BARRY\_VERSION\_MAYOR ## . ## BARRY\_VERSION\_MINOR
- #define COUNTER\_FUNCTION(a)
- #define COUNTER\_LAMBDA(a)
- #define RULE\_FUNCTION(a)
- #define RULE\_LAMBDA(a)

#### 9.21.1 Macro Definition Documentation

#### 9.21.1.1 BARRY HPP

```
#define BARRY_HPP
```

Definition at line 29 of file barry.hpp.

#### 9.21.1.2 BARRY\_VERSION

```
#define BARRY_VERSION BARRY_VERSION_MAYOR ## . ## BARRY_VERSION_MINOR
```

Definition at line 33 of file barry.hpp.

#### 9.21.1.3 BARRY\_VERSION\_MAYOR

```
#define BARRY_VERSION_MAYOR 0
```

Definition at line 31 of file barry.hpp.

## 9.21.1.4 BARRY\_VERSION\_MINOR

```
#define BARRY_VERSION_MINOR 1
```

Definition at line 32 of file barry.hpp.

## 9.21.1.5 COUNTER\_FUNCTION

```
\begin{array}{c} \texttt{\#define COUNTER\_FUNCTION}\,(\\ & a \end{array})
```

#### Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
inline double (a) (const Array_Type & Array, size_t i, size_t j, Data_Type & data)\
```

Definition at line 92 of file barry.hpp.

## 9.21.1.6 COUNTER\_LAMBDA

Definition at line 95 of file barry.hpp.

### 9.21.1.7 RULE\_FUNCTION

Definition at line 99 of file barry.hpp.

#### 9.21.1.8 **RULE LAMBDA**

Definition at line 102 of file barry.hpp.

# 9.22 include/barry/cell-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Classes**

class Cell
 Cell\_Type >
 Entries in BArray. For now, it only has two members:

# 9.23 include/barry/cell-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



# 9.24 include/barry/col-bones.hpp File Reference

# 9.25 include/barry/counters-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:

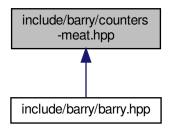


### **Classes**

- class Counter< Array\_Type, Data\_Type >
  - A counter function based on change statistics.

## 9.26 include/barry/counters-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Macros**

- #define COUNTER\_TYPE() Counter<Array\_Type,Data\_Type>
- #define COUNTER\_TEMPLATE\_ARGS() < typename Array\_Type, typename Data\_Type>
- #define COUNTER\_TEMPLATE(a, b) template COUNTER\_TEMPLATE\_ARGS() inline a COUNTER\_TYPE() 

  ::b
- #define TMP\_HASHER\_CALL Hasher\_fun\_type<Array\_Type,Data\_Type>
- #define COUNTERS\_TYPE() Counters<Array\_Type,Data\_Type>
- #define COUNTERS\_TEMPLATE\_ARGS() < typename Array\_Type, typename Data\_Type>
- #define COUNTERS\_TEMPLATE(a, b) template COUNTERS\_TEMPLATE\_ARGS() inline a COUNTERS\_TYPE()

   ::b

## **Functions**

- COUNTER\_TEMPLATE (, Counter)(const Counter< Array Type
- Data\_Type init\_fun (counter\_.init\_fun)
- Data\_Type hasher\_fun (counter\_.hasher\_fun)
- Data\_Type &&counter\_\_init\_fun (std::move(counter\_\_init\_fun))
- Data\_Type &&counter\_ hasher\_fun (std::move(counter\_.hasher\_fun))
- Data\_Type &&counter\_ data (std::move(counter\_.data))
- Data Type &&counter name (std::move(counter .name))
- Data\_Type &&counter\_ desc (std::move(counter\_.desc))

#### Move constructor.

- COUNTER\_TEMPLATE (COUNTER\_TYPE(), operator=)(const Counter< Array\_Type
- COUNTER\_TEMPLATE (COUNTER\_TYPE() &, operator=)(Counter< Array\_Type
- COUNTER\_TEMPLATE (double, count)(Array\_Type &Array

#### < Move assignment

- return count\_fun (Array, i, j, data)
- COUNTER TEMPLATE (double, init)(Array Type & Array
- return init\_fun (Array, i, j, data)
- COUNTER\_TEMPLATE (std::string, get\_name)() const
- COUNTER\_TEMPLATE (std::string, get\_description)() const
- COUNTER\_TEMPLATE (void, set\_hasher)(Hasher\_fun\_type< Array\_Type</li>

- COUNTER\_TEMPLATE (TMP\_HASHER\_CALL, get\_hasher)()
- COUNTERS\_TEMPLATE (, Counters)()
- COUNTERS\_TEMPLATE (COUNTER\_TYPE() &, operator[])(size\_t idx)
- Data\_Type hasher (counter\_.hasher)
- Data Type &&counters hasher (std::move(counters .hasher))
- COUNTERS TEMPLATE (COUNTERS TYPE(), operator=)(const Counters < Array Type</li>
- COUNTERS\_TEMPLATE (COUNTERS\_TYPE() &, operator=)(Counters< Array\_Type
- COUNTERS TEMPLATE (void, add counter)(Counter< Array Type</li>
- COUNTERS\_TEMPLATE (std::vector< std::string >, get\_names)() const
- COUNTERS\_TEMPLATE (std::vector< std::string >, get\_descriptions)() const
- COUNTERS TEMPLATE (std::vector< double >, gen hash)(const Array Type & array
- for (auto &c:data)
- if (add\_dims)
- if (hasher)
- if (res.size()==0u) res.push\_back(0.0)
- COUNTERS\_TEMPLATE (void, add\_hash)(Hasher\_fun\_type< Array\_Type</li>

#### **Variables**

- Data Type & counter
- Data Type &&counter noexcept
- size ti
- size\_t size\_t j
- Data\_Type fun
- Data Type counter
- · return
- Data\_Type count\_fun\_
- Data\_Type Counter\_fun\_type< Array\_Type, Data\_Type > init\_fun\_
- Data\_Type Counter\_fun\_type
   Array\_Type, Data\_Type > Hasher\_fun\_type
   Array\_Type, Data\_Type > hasher\_fun\_
- Data\_Type Counter\_fun\_type
   Array\_Type, Data\_Type > Hasher\_fun\_type
   Array\_Type, Data\_Type > Data\_Type data\_
- Data\_Type Counter\_fun\_type
   Array\_Type, Data\_Type > Hasher\_fun\_type
   Array\_Type, Data\_Type > Data\_Type std::string name
- Data\_Type Counter\_fun\_type
   Array\_Type, Data\_Type > Hasher\_fun\_type
   Array\_Type, Data\_Type > Data\_Type std::string std::string desc\_
- · bool add dims
- · return res
- Data\_Type fun\_

#### 9.26.1 Macro Definition Documentation

#### 9.26.1.1 COUNTER\_TEMPLATE

Definition at line 8 of file counters-meat.hpp.

#### 9.26.1.2 COUNTER\_TEMPLATE\_ARGS

```
#define COUNTER_TEMPLATE_ARGS() <typename Array_Type, typename Data_Type>
```

Definition at line 6 of file counters-meat.hpp.

#### 9.26.1.3 COUNTER\_TYPE

```
#define COUNTER_TYPE( ) Counter<Array_Type,Data_Type>
```

Definition at line 4 of file counters-meat.hpp.

#### 9.26.1.4 COUNTERS\_TEMPLATE

Definition at line 129 of file counters-meat.hpp.

#### 9.26.1.5 COUNTERS\_TEMPLATE\_ARGS

```
#define COUNTERS_TEMPLATE_ARGS() <typename Array_Type, typename Data_Type>
```

Definition at line 127 of file counters-meat.hpp.

### 9.26.1.6 COUNTERS\_TYPE

```
#define COUNTERS_TYPE( ) Counters<Array_Type,Data_Type>
```

Definition at line 125 of file counters-meat.hpp.

## 9.26.1.7 TMP\_HASHER\_CALL

```
#define TMP_HASHER_CALL Hasher_fun_type<Array_Type,Data_Type>
```

Definition at line 115 of file counters-meat.hpp.

#### 9.26.2 Function Documentation

#### 9.26.2.1 count fun()

## 9.26.2.2 COUNTER\_TEMPLATE() [1/9]

```
COUNTER_TEMPLATE (

Counter ) const
```

## 9.26.2.3 COUNTER\_TEMPLATE() [2/9]

## 9.26.2.4 COUNTER\_TEMPLATE() [3/9]

## 9.26.2.5 COUNTER\_TEMPLATE() [4/9]

< Move assignment

## **9.26.2.6 COUNTER\_TEMPLATE()** [5/9]

## 9.26.2.7 **COUNTER\_TEMPLATE()** [6/9]

```
COUNTER_TEMPLATE (
          std::string ,
          get_description ) const
```

Definition at line 107 of file counters-meat.hpp.

## 9.26.2.8 **COUNTER\_TEMPLATE()** [7/9]

```
COUNTER_TEMPLATE (
          std::string ,
          get_name ) const
```

Definition at line 103 of file counters-meat.hpp.

### 9.26.2.9 COUNTER\_TEMPLATE() [8/9]

```
COUNTER_TEMPLATE (

TMP_HASHER_CALL ,

qet_hasher )
```

Definition at line 116 of file counters-meat.hpp.

#### 9.26.2.10 COUNTER\_TEMPLATE() [9/9]

## 9.26.2.11 COUNTERS\_TEMPLATE() [1/9]

```
COUNTERS_TEMPLATE (

Counters )
```

Definition at line 132 of file counters-meat.hpp.

#### 9.26.2.12 COUNTERS\_TEMPLATE() [2/9]

```
COUNTERS_TEMPLATE (

COUNTER_TYPE() & ,

operator [])
```

Definition at line 134 of file counters-meat.hpp.

## 9.26.2.13 COUNTERS\_TEMPLATE() [3/9]

## 9.26.2.14 COUNTERS\_TEMPLATE() [4/9]

```
COUNTERS_TEMPLATE (

COUNTERS_TYPE() ,

operator ) const
```

## 9.26.2.15 COUNTERS\_TEMPLATE() [5/9]

```
COUNTERS_TEMPLATE (
          std::vector< double > ,
          gen_hash ) const &
```

#### 9.26.2.16 COUNTERS\_TEMPLATE() [6/9]

```
COUNTERS_TEMPLATE (
          std::vector< std::string > ,
          get_descriptions ) const
```

Definition at line 213 of file counters-meat.hpp.

## 9.26.2.17 COUNTERS\_TEMPLATE() [7/9]

```
COUNTERS_TEMPLATE (
          std::vector< std::string > ,
          get_names ) const
```

Definition at line 201 of file counters-meat.hpp.

## 9.26.2.18 COUNTERS\_TEMPLATE() [8/9]

## 9.26.2.19 **COUNTERS\_TEMPLATE()** [9/9]

## 9.26.2.20 data()

## 9.26.2.21 desc()

Move constructor.

Definition at line 32 of file counters-meat.hpp.

### 9.26.2.22 for()

```
for ( auto &c:data )
```

Definition at line 233 of file counters-meat.hpp.

## 9.26.2.23 hasher() [1/2]

Definition at line 141 of file counters-meat.hpp.

## 9.26.2.24 hasher() [2/2]

Definition at line 144 of file counters-meat.hpp.

## 9.26.2.25 hasher\_fun() [1/2]

Definition at line 13 of file counters-meat.hpp.

#### 9.26.2.26 hasher\_fun() [2/2]

## 9.26.2.27 if() [1/3]

```
if (
    add_dims )
```

Definition at line 248 of file counters-meat.hpp.

## 9.26.2.28 if() [2/3]

```
if ( hasher )
```

Definition at line 255 of file counters-meat.hpp.

```
9.26.2.29 if() [3/3]
```

```
if ( {\tt res.} \quad {\tt size() = =0u \ )}
```

### 9.26.2.30 init\_fun() [1/3]

## 9.26.2.31 init\_fun() [2/3]

## 9.26.2.32 init\_fun() [3/3]

### 9.26.2.33 name()

## 9.26.3 Variable Documentation

#### 9.26.3.1 add\_dims

```
bool add_dims
```

#### Initial value:

```
std::vector<double> res
```

Definition at line 227 of file counters-meat.hpp.

#### 9.26.3.2 count\_fun\_

```
Data_Type count_fun_
```

Definition at line 179 of file counters-meat.hpp.

#### 9.26.3.3 counter

```
Data_Type counter

Initial value:
{
    data.push_back(counter)
```

Definition at line 170 of file counters-meat.hpp.

## 9.26.3.4 counter\_

```
Data_Type & counter_

Initial value:
{
    if (this != &counter_) {
        this->count_fun = counter_.count_fun;
        this->init_fun = counter_.init_fun;
        this->hasher_fun = counter_.hasher_fun;

        this->data = counter_.data;
        this->name = counter_.name;
        this->desc = counter_.desc;
    }
    return *this
```

Definition at line 12 of file counters-meat.hpp.

## 9.26.3.5 data\_

Data\_Type Counter\_fun\_type<Array\_Type,Data\_Type> Hasher\_fun\_type<Array\_Type,Data\_Type> Data←
\_Type data\_

Definition at line 182 of file counters-meat.hpp.

### 9.26.3.6 desc\_

```
Data_Type Counter_fun_type<Array_Type,Data_Type> Hasher_fun_type<Array_Type,Data_Type> Data←
_Type std::string std::string desc_
```

### Initial value:

```
data.emplace_back(Counter<Array_Type,Data_Type>(
    count_fun_,
    init_fun_,
    hasher_fun_,
    data_,
    name_,
    desc_
))
```

Definition at line 184 of file counters-meat.hpp.

#### 9.26.3.7 fun

```
Data_Type fun

Initial value:
{
    hasher_fun = fun
```

Definition at line 111 of file counters-meat.hpp.

### 9.26.3.8 fun

```
Data_Type fun_
```

# Initial value:

```
hasher = fun_
```

Definition at line 270 of file counters-meat.hpp.

# 9.26.3.9 hasher\_fun\_

Data\_Type Counter\_fun\_type<Array\_Type,Data\_Type> Hasher\_fun\_type<Array\_Type,Data\_Type> hasher← \_fun\_

Definition at line 181 of file counters-meat.hpp.

### 9.26.3.10 i

```
size_t i
```

Definition at line 83 of file counters-meat.hpp.

### 9.26.3.11 init\_fun\_

```
Data_Type Counter_fun_type<Array_Type,Data_Type> init_fun_
```

Definition at line 180 of file counters-meat.hpp.

### 9.26.3.12 j

```
size_t j

Initial value:
{
    if (count_fun == nullptr)
        return 0.0
```

Definition at line 83 of file counters-meat.hpp.

### 9.26.3.13 name\_

```
Data_Type Counter_fun_type<Array_Type,Data_Type> Hasher_fun_type<Array_Type,Data_Type> Data←
_Type std::string name_
```

Definition at line 183 of file counters-meat.hpp.

### 9.26.3.14 noexcept

```
Data_Type &&counters_ noexcept

Initial value:
{
    if (this != &counter_)
    {
        this->data = std::move(counter_.data);

        this->count_fun = std::move(counter_.count_fun);
        this->init_fun = std::move(counter_.init_fun);
        this->hasher_fun = std::move(counter_.hasher_fun);

        this->name = std::move(counter_.name);
        this->desc = std::move(counter_.desc);
    }
    return *this
```

Definition at line 26 of file counters-meat.hpp.

#### 9.26.3.15 res

return res

Definition at line 265 of file counters-meat.hpp.

### 9.26.3.16 return

return

Definition at line 175 of file counters-meat.hpp.

# 9.27 include/barry/counters/network-css.hpp File Reference

This graph shows which files directly or indirectly include this file:



### **Macros**

- #define CSS\_SIZE()
- #define CSS\_MATCH\_TYPE()
- #define CSS\_CASE\_TRUTH() if ((i < n) && (j < n))
- #define CSS\_TRUE\_CELLS()
- #define CSS\_CASE\_PERCEIVED() else if (((i >= s) && (i < e)) & ((j >= s) && (j < e)))
- #define CSS\_PERCEIVED\_CELLS()
- #define CSS\_CASE\_ELSE()
- #define CSS\_CHECK\_SIZE\_INIT()
- #define CSS\_CHECK\_SIZE()
- #define CSS\_APPEND(name)
- #define CSS\_NET\_COUNTER\_LAMBDA\_INIT()

#### **Functions**

```
• template<typename Tnet = Network>
  void counter css partially false recip commi (NetCounters< Tnet > *counters, size t netsize, const std ←
  ::vector< size_t > &end_, size_t counter_type=0u)
     Counts errors of commission.
• template<typename Tnet = Network>
  void counter css partially false recip omiss (NetCounters< Tnet > *counters, size t netsize, const std↔
  ::vector< size t > &end , size t counter type=0u)
     Counts errors of omission.
• template<typename Tnet = Network>
  void counter css completely false recip comiss (NetCounters< Tnet > *counters, size t netsize, const
  std::vector< size_t > &end_, size_t counter_type=0u)
     Counts completely false reciprocity (comission)
template<typename Tnet = Network>
  void counter css completely false recip omiss (NetCounters< Tnet > *counters, size t netsize, const
  std::vector< size_t > &end_, size_t counter_type=0u)
     Counts completely false reciprocity (omission)
template<typename Tnet = Network>
  void counter css mixed recip (NetCounters< Tnet > *counters, size t netsize, const std::vector< size t >
  &end , size t counter type=0u)
     Counts mixed reciprocity errors.
• template<typename Tnet = Network>
  void counter_css_census01 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end_, size_t counter_type=0u)

    template<tvpename Tnet = Network>

  void counter_css_census02 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end , size t counter type=0u)

    template<typename Tnet = Network>

  void counter_css_census03 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end_, size_t counter_type=0u)
template<typename Tnet = Network>
  void counter_css_census04 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end_, size_t counter_type=0u)
template<typename Tnet = Network>
  void counter_css_census05 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end_, size_t counter_type=0u)
template<typename Tnet = Network>
  void counter css census06 (NetCounters< Tnet > *counters, size t netsize, const std::vector< size t >
  &end_, size_t counter_type=0u)
template<typename Tnet = Network>
  void counter_css_census07 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end_, size_t counter_type=0u)
template<typename Tnet = Network>
  void counter_css_census08 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end , size t counter type=0u)
template<typename Tnet = Network>
  void counter css census09 (NetCounters< Tnet > *counters, size t netsize, const std::vector< size t >
  &end_, size_t counter_type=0u)
template<typename Tnet = Network>
  void counter css census10 (NetCounters< Tnet > *counters, size t netsize, const std::vector< size t >
  &end , size t counter type=0u)
```

### 9.27.1 Macro Definition Documentation

### 9.27.1.1 CSS\_APPEND

Definition at line 81 of file network-css.hpp.

### 9.27.1.2 CSS\_CASE\_ELSE

```
#define CSS_CASE_ELSE( )
```

Definition at line 66 of file network-css.hpp.

### 9.27.1.3 CSS CASE PERCEIVED

```
      \# define \ CSS\_CASE\_PERCEIVED ( ) \ else \ if \ (((i >= s) \ \&\& \ (i < e)) \ \& \ ((j >= s) \ \&\& \ (j < e)))
```

Definition at line 48 of file network-css.hpp.

### 9.27.1.4 CSS\_CASE\_TRUTH

```
#define CSS_CASE_TRUTH( ) if ((i < n) && (j < n))
```

Definition at line 32 of file network-css.hpp.

### 9.27.1.5 CSS CHECK SIZE

Definition at line 76 of file network-css.hpp.

### 9.27.1.6 CSS\_CHECK\_SIZE\_INIT

Definition at line 70 of file network-css.hpp.

### 9.27.1.7 CSS\_MATCH\_TYPE

Definition at line 22 of file network-css.hpp.

## 9.27.1.8 CSS\_NET\_COUNTER\_LAMBDA\_INIT

Definition at line 89 of file network-css.hpp.

### 9.27.1.9 CSS PERCEIVED CELLS

```
#define CSS_PERCEIVED_CELLS()

Value:
    size_t i_ = i - s; \
    size_t j_ = j - s; \
    CSS_MATCH_TYPE() \
    double tji = static_cast<double>(Array(j - s, i - s, false)); \
    double pji = static_cast<double>(Array(j, i, false)); \
    double tij = static_cast<double>(Array(i - s, j - s, false));
```

Definition at line 55 of file network-css.hpp.

### 9.27.1.10 CSS\_SIZE

```
#define CSS_SIZE()

Value:
    size_t n = data.indices[0u]; \
    size_t s = data.indices[1u]; \
    size_t e = data.indices[2u]; \
    size_t ctype = data.indices[3u]; \
    size_t ego_id = data.indices[4u]; \
    if (ctype > 2) \
        throw std::range_error("Counter type should be 0, 1, or 2.");
```

Definition at line 8 of file network-css.hpp.

### 9.27.1.11 CSS\_TRUE\_CELLS

```
#define CSS_TRUE_CELLS( )

Value:
    size_t i_ = i; \
    size_t j_ = j; \
    CSS_MATCH_TYPE() \
    double tji = static_cast<double>(Array(j, i, false)); \
    double pij = static_cast<double>(Array(i + s, j + s, false)); \
    double pji = static_cast<double>(Array(j + s, i + s, false)); \
```

Definition at line 39 of file network-css.hpp.

### 9.27.2 Function Documentation

### 9.27.2.1 counter css census01()

Definition at line 324 of file network-css.hpp.

# 9.27.2.2 counter\_css\_census02()

Definition at line 389 of file network-css.hpp.

### 9.27.2.3 counter\_css\_census03()

Definition at line 429 of file network-css.hpp.

#### 9.27.2.4 counter css census04()

Definition at line 469 of file network-css.hpp.

### 9.27.2.5 counter\_css\_census05()

Definition at line 509 of file network-css.hpp.

### 9.27.2.6 counter\_css\_census06()

Definition at line 549 of file network-css.hpp.

### 9.27.2.7 counter\_css\_census07()

Definition at line 589 of file network-css.hpp.

#### 9.27.2.8 counter css census08()

Definition at line 629 of file network-css.hpp.

### 9.27.2.9 counter\_css\_census09()

Definition at line 669 of file network-css.hpp.

### 9.27.2.10 counter\_css\_census10()

Definition at line 709 of file network-css.hpp.

### 9.27.2.11 counter\_css\_completely\_false\_recip\_comiss()

Counts completely false reciprocity (comission)

Definition at line 200 of file network-css.hpp.

### 9.27.2.12 counter\_css\_completely\_false\_recip\_omiss()

Counts completely false reciprocity (omission)

Definition at line 241 of file network-css.hpp.

# 9.27.2.13 counter\_css\_mixed\_recip()

Counts mixed reciprocity errors.

Definition at line 282 of file network-css.hpp.

### 9.27.2.14 counter\_css\_partially\_false\_recip\_commi()

Counts errors of commission.

#### **Parameters**

netsize	Size of the reference (true) network
end_	Vector indicating one past the ending index of each network. (see details)
counter_type	Size_t indicating the type of counter to use. Possible values are: 0: Count all, 1: Only count if perceiver is involved, and 2: Only count if perceiver is not involved.

The  ${\tt end\_parameter}$  should be of length N of  ${\tt networks-1}$ . It is assumed that the first network ends at  ${\tt netsize}$ .

Definition at line 107 of file network-css.hpp.

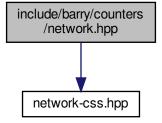
# 9.27.2.15 counter\_css\_partially\_false\_recip\_omiss()

Counts errors of omission.

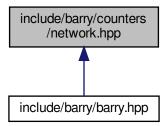
Definition at line 155 of file network-css.hpp.

# 9.28 include/barry/counters/network.hpp File Reference

#include "network-css.hpp"
Include dependency graph for network.hpp:



This graph shows which files directly or indirectly include this file:



### **Classes**

class NetworkData

Data class for Networks.

· class NetCounterData

Data class used to store arbitrary size\_t or double vectors.

### **Macros**

- #define NET\_C\_DATA\_IDX(i) (data.indices[i])
- #define NET\_C\_DATA\_NUM(i) (data.numbers[i])

# Macros for defining counters

- #define NETWORK\_COUNTER(a)
- #define NETWORK\_COUNTER\_LAMBDA(a)
- #define NETWORKDENSE\_COUNTER\_LAMBDA(a)

# Macros for defining rules

- #define NETWORK\_RULE(a)
- #define NETWORK\_RULE\_LAMBDA(a)

#### **Functions**

```
• template<typename Tnet = Network>
  void counter_edges (NetCounters < Tnet > *counters)
     Number of edges.
template<typename Tnet = Network>
  void counter isolates (NetCounters< Tnet > *counters)
     Number of isolated vertices.

    template<> void counter_isolates (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter mutual (NetCounters < Tnet > *counters)
     Number of mutual ties.
• template<typename Tnet = Network>
  void counter_istar2 (NetCounters< Tnet > *counters)

    template<> void counter istar2 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter ostar2 (NetCounters< Tnet > *counters)

    template<> void counter ostar2 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter ttriads (NetCounters< Tnet > *counters)

    template<> void counter_ttriads (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter_ctriads (NetCounters< Tnet > *counters)

    template<> void counter_ctriads (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter_density (NetCounters < Tnet > *counters)
• template<typename Tnet = Network>
  void counter_idegree15 (NetCounters< Tnet > *counters)

    template<> void counter idegree15 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter odegree15 (NetCounters< Tnet > *counters)

    template<> void counter_odegree15 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter_absdiff (NetCounters < Tnet > *counters, size_t attr_id, double alpha=1.0)
     Sum of absolute attribute difference between ego and alter.
• template<typename Tnet = Network>
  void counter diff (NetCounters < Tnet > *counters, size t attr id, double alpha=1.0, double tail head=true)
     Sum of attribute difference between ego and alter to pow(alpha)

    NETWORK_COUNTER (init_single_attr)

    template<typename Tnet = Network>

  void counter nodeicov (NetCounters < Tnet > *counters, size t attr id)
template<typename Tnet = Network>
  void counter nodeocov (NetCounters< Tnet > *counters, size t attr id)
template<typename Tnet = Network>
  void counter_nodecov (NetCounters< Tnet > *counters, size_t attr_id)
template<typename Tnet = Network>
  void counter_nodematch (NetCounters < Tnet > *counters, size_t attr_id)
• template<typename Tnet = Network>
  void counter_idegree (NetCounters< Tnet > *counters, std::vector< size_t > d)
     Counts number of vertices with a given in-degree.

    template<> void counter_idegree (NetCounters< NetworkDense > *counters, std::vector< size_t > d)

template<typename Tnet = Network>
  void counter_odegree (NetCounters< Tnet > *counters, std::vector< size_t > d)
     Counts number of vertices with a given out-degree.

    template<> void counter_odegree (NetCounters< NetworkDense > *counters, std::vector< size_t > d)
```

```
    template < typename Tnet = Network>
    void counter_degree (NetCounters < Tnet > *counters, std::vector < size_t > d)
        Counts number of vertices with a given out-degree.
```

### Rules for network models

#### **Parameters**

```
rules | A pointer to a NetRules object (Rules < Network, bool > ).
```

template<typename Tnet = Network>
 void rules\_zerodiag (NetRules< Tnet > \*rules)
 Number of edges.

# Convenient typedefs for network objects.

- #define BARRY ZERO NETWORK 0.0
- #define BARRY\_ZERO\_NETWORK\_DENSE 0
- typedef BArray< double, NetworkData > Network
- typedef BArrayDense< int, NetworkData > NetworkDense
- template < typename Tnet = Network > using NetCounter = Counter < Tnet, NetCounterData >
- template < typename Tnet = Network >
   using NetCounters = Counters < Tnet, NetCounterData >
- template<typename Tnet = Network>
   using NetSupport = Support< Tnet, NetCounterData >
- template<typename Tnet = Network>
   using NetStatsCounter = StatsCounter< Tnet, NetCounterData >
- template < typename Tnet >
   using NetModel = Model < Tnet, NetCounterData >
- template<typename Tnet = Network>
   using NetRule = Rule< Tnet, bool >
- template<typename Tnet = Network>
   using NetRules = Rules< Tnet, bool >

### 9.28.1 Macro Definition Documentation

### 9.28.1.1 BARRY\_ZERO\_NETWORK

#define BARRY ZERO NETWORK 0.0

Definition at line 85 of file network.hpp.

# 9.28.1.2 BARRY\_ZERO\_NETWORK\_DENSE

```
#define BARRY_ZERO_NETWORK_DENSE 0
```

Definition at line 86 of file network.hpp.

# 9.28.1.3 NET\_C\_DATA\_IDX

Definition at line 74 of file network.hpp.

### 9.28.1.4 NET\_C\_DATA\_NUM

Definition at line 75 of file network.hpp.

### 9.28.1.5 NETWORK\_COUNTER

#### Value:

```
template<typename Tnet = Network>\
inline double (a) (const Tnet & Array, size_t i, size_t j, NetCounterData & data)
```

Function for definition of a network counter function

Definition at line 114 of file network.hpp.

### 9.28.1.6 NETWORK\_COUNTER\_LAMBDA

```
#define NETWORK_COUNTER_LAMBDA( a )
```

### Value:

```
Counter_fun_type<Tnet, NetCounterData> a = \
   [](const Tnet & Array, size_t i, size_t j, NetCounterData & data)
```

Lambda function for definition of a network counter function

Definition at line 119 of file network.hpp.

# 9.28.1.7 NETWORK\_RULE

#### Value:

```
template<typename Tnet = Network>\
inline bool (a) (const Tnet & Array, size_t i, size_t j, bool & data)
```

Function for definition of a network counter function

Definition at line 133 of file network.hpp.

### 9.28.1.8 NETWORK RULE LAMBDA

Lambda function for definition of a network counter function

Definition at line 138 of file network.hpp.

### 9.28.1.9 NETWORKDENSE\_COUNTER\_LAMBDA

```
Counter_fun_type<NetworkDense, NetCounterData> a = \
   [](const NetworkDense & Array, size_t i, size_t j, NetCounterData & data)
```

Definition at line 123 of file network.hpp.

# 9.28.2 Typedef Documentation

### 9.28.2.1 NetCounter

```
template<typename Tnet = Network>
using NetCounter = Counter<Tnet, NetCounterData >
```

Definition at line 89 of file network.hpp.

### 9.28.2.2 NetCounters

```
template<typename Tnet = Network>
using NetCounters = Counters<Tnet, NetCounterData>
```

Definition at line 92 of file network.hpp.

### 9.28.2.3 NetModel

```
template<typename Tnet >
using NetModel = Model<Tnet, NetCounterData>
```

Definition at line 101 of file network.hpp.

### 9.28.2.4 NetRule

```
template<typename Tnet = Network>
using NetRule = Rule<Tnet, bool>
```

Definition at line 104 of file network.hpp.

### 9.28.2.5 NetRules

```
template<typename Tnet = Network>
using NetRules = Rules<Tnet, bool>
```

Definition at line 107 of file network.hpp.

#### 9.28.2.6 NetStatsCounter

```
template<typename Tnet = Network>
using NetStatsCounter = StatsCounter<Tnet, NetCounterData>
```

Definition at line 98 of file network.hpp.

# 9.28.2.7 NetSupport

```
template<typename Tnet = Network>
using NetSupport = Support<Tnet, NetCounterData >
```

Definition at line 95 of file network.hpp.

### 9.28.2.8 Network

```
typedef BArray<double, NetworkData> Network
```

Definition at line 82 of file network.hpp.

### 9.28.2.9 NetworkDense

```
typedef BArrayDense<int, NetworkData> NetworkDense
```

Definition at line 83 of file network.hpp.

# 9.28.3 Function Documentation

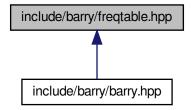
# 9.28.3.1 rules\_zerodiag()

Number of edges.

Definition at line 1381 of file network.hpp.

# 9.29 include/barry/freqtable.hpp File Reference

This graph shows which files directly or indirectly include this file:



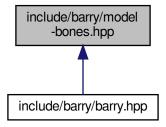
### **Classes**

class FreqTable
 T >

Frequency table of vectors.

# 9.30 include/barry/model-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



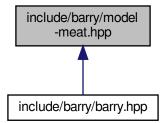
# Classes

- class Model < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type >  $\,$ 

General framework for discrete exponential models. This class allows generating discrete exponential models in the form of a linear exponential model:

# 9.31 include/barry/model-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



### **Functions**

- double update\_normalizing\_constant (const std::vector< double > &params, const double \*support, size\_t k, size t n)
- double likelihood\_ (const double \*stats\_target, const std::vector< double > &params, const double normalizing\_constant, size\_t n\_params, bool log\_=false)

### 9.31.1 Function Documentation

### 9.31.1.1 likelihood\_()

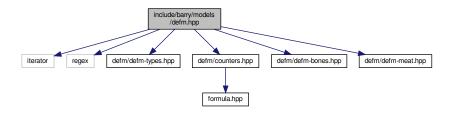
Definition at line 71 of file model-meat.hpp.

### 9.31.1.2 update\_normalizing\_constant()

Definition at line 9 of file model-meat.hpp.

# 9.32 include/barry/models/defm.hpp File Reference

```
#include <iterator>
#include <regex>
#include "defm/defm-types.hpp"
#include "defm/counters.hpp"
#include "defm/defm-bones.hpp"
#include "defm/defm-meat.hpp"
Include dependency graph for defm.hpp:
```

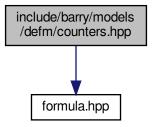


# **Namespaces**

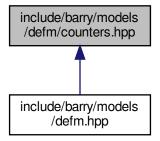
• defm

# 9.33 include/barry/models/defm/counters.hpp File Reference

#include "formula.hpp"
Include dependency graph for counters.hpp:



This graph shows which files directly or indirectly include this file:



### **Macros**

- #define MAKE\_DEFM\_HASHER(hasher, a, cov)
- #define DEFM\_RULEDYN\_LAMBDA(a)
- #define UNI\_SUB(a)

### Macros for defining counters

- #define DEFM\_COUNTER(a) inline double (a) (const DEFMArray & Array, size\_t i, size\_t j, DEFMCounterData & data)
- #define DEFM\_COUNTER\_LAMBDA(a)

#### Macros for defining rules

- #define DEFM\_RULE(a) inline bool (a) (const DEFMArray & Array, size\_t i, size\_t j, bool & data)
- #define DEFM\_RULE\_LAMBDA(a)

### **Functions**

void counter\_ones (DEFMCounters \*counters, int covar\_index=-1, std::string vname="", const std::vector< std::string > \*x\_names=nullptr)

Prevalence of ones.

- void counter\_logit\_intercept (DEFMCounters \*counters, size\_t n\_y, std::vector< size\_t > which={}, int covar\_index=-1, std::string vname="", const std::vector< std::string > \*x\_names=nullptr, const std::vector< std::string > \*y\_names=nullptr)

Prevalence of ones.

void counter\_transition\_formula (DEFMCounters \*counters, std::string formula, size\_t m\_order, size\_t n\_y, int covar\_index=-1, std::string vname="", const std::vector< std::string > \*x\_names=nullptr, const std::vector< std::string > \*y\_names=nullptr)

Prevalence of ones.

• void counter\_fixed\_effect (DEFMCounters \*counters, int covar\_index, double k, std::string vname="", const std::vector< std::string > \*x names=nullptr)

Prevalence of ones.

### Returns true if the cell is free

### **Parameters**

rules | A pointer to a DEFMRules object (Rules < DEFMArray, bool > ).

- void rules\_markov\_fixed (DEFMRules \*rules, size\_t markov\_order)
  - Number of edges.
- void rules\_dont\_become\_zero (DEFMSupport \*support, std::vector< size\_t > ids)

Blocks switching a one to zero.

### 9.33.1 Macro Definition Documentation

# 9.33.1.1 DEFM\_COUNTER

Function for definition of a network counter function

Definition at line 39 of file counters.hpp.

### 9.33.1.2 DEFM\_COUNTER\_LAMBDA

[](const DEFMArray & Array, size\_t i, size\_t j, DEFMCounterData & data) -> double

Lambda function for definition of a network counter function

barry::Counter\_fun\_type<DEFMArray, DEFMCounterData> a = \

Definition at line 43 of file counters.hpp.

# 9.33.1.3 DEFM\_RULE

Function for definition of a network counter function

Definition at line 53 of file counters.hpp.

# 9.33.1.4 DEFM\_RULE\_LAMBDA

#### Value:

```
barry::Rule_fun_type<DEFMArray, DEFMRuleData> a = \
[](const DEFMArray & Array, size_t i, size_t j, DEFMRuleData & data) -> bool
```

Lambda function for definition of a network counter function

Definition at line 57 of file counters.hpp.

### 9.33.1.5 DEFM\_RULEDYN\_LAMBDA

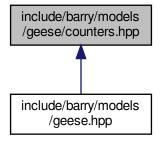
Lambda function for definition of a network counter function

Definition at line 63 of file counters.hpp.

# 9.33.1.6 UNI\_SUB

# 9.34 include/barry/models/geese/counters.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Functions**

- void rule\_leafs (PhyloSupport \*support)
- void rule\_dyn\_limit\_changes (PhyloSupport \*support, size\_t pos, size\_t lb, size\_t ub, size\_
   t duplication=Geese::etype\_default)

Overall functional gains.

- #define MAKE DUPL VARS()
- #define IS\_EITHER() (DATA\_AT == Geese::etype\_either)
- #define IS\_DUPLICATION() ((DATA\_AT == Geese::etype\_duplication) & (DPL))
- #define IS\_SPECIATION() ((DATA\_AT == Geese::etype\_speciation) & (!DPL))
- #define IF\_MATCHES()
- #define IF\_NOTMATCHES()
- #define PHYLO\_RULE\_LAMBDA(a)

Extension of a simple counter.

- #define PHYLO COUNTER LAMBDA(a)
- #define PHYLO\_RULE\_DYN\_LAMBDA(a)
- #define PHYLO CHECK MISSING()
- std::string get last name (size t d)
- void counter\_overall\_gains (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

Overall functional gains.

- void counter\_gains (PhyloCounters \*counters, std::vector < size\_t > nfun, size\_t duplication=Geese::etype\_default)

  Functional gains for a specific function (nfun).
- void counter\_gains\_k\_offspring (PhyloCounters \*counters, std::vector< size\_t > nfun, size\_t k=1u, size\_t duplication=Geese::etype\_default)

k genes gain function nfun

void counter genes changing (PhyloCounters \*counters, size t duplication=Geese::etype default)

Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)

• void counter\_preserve\_pseudogene (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_
t duplication=Geese::etype\_default)

Keeps track of how many pairs of genes preserve pseudostate.

• void counter prop genes changing (PhyloCounters \*counters, size t duplication=Geese::etype default)

Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)

void counter\_overall\_loss (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

Overall functional loss.

- void counter\_maxfuns (PhyloCounters \*counters, size\_t lb, size\_t ub, size\_t duplication=Geese::etype\_default)

  Cap the number of functions per gene.
- void counter\_loss (PhyloCounters \*counters, std::vector < size\_t > nfun, size\_t duplication=Geese::etype\_default)

  Total count of losses for an specific function.
- void counter\_overall\_changes (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

Total number of changes. Use this statistic to account for "preservation".

- void counter\_subfun (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

  Total count of Sub-functionalization events.
- void counter\_cogain (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

  Co-evolution (joint gain or loss)
- void counter\_longest (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

Longest branch mutates (either by gain or by loss)

- void counter\_neofun (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

  Total number of neofunctionalization events.
- void counter\_pairwise\_neofun\_singlefun (PhyloCounters \*counters, size\_t nfunA, size\_t duplication=Geese::etype\_default)
   Total number of neofunctionalization events sum\_u sum\_{{w < u}} [x(u,a)\*(1 x(w,a)) + (1 x(u,a)) \* x(w,a)] change stat: delta{x(u,a): 0->1} = 1 2 \* x(w,a)

• void counter\_neofun\_a2b (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

Total number of neofunctionalization events.

- void counter\_co\_opt (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default) Function co-opting.
- void counter\_k\_genes\_changing (PhyloCounters \*counters, size\_t k, size\_t duplication=Geese::etype\_default)

  Indicator function. Equals to one if k genes changed and zero otherwise.
- void counter\_less\_than\_p\_prop\_genes\_changing (PhyloCounters \*counters, double p, size\_t duplication=Geese::etype\_default Indicator function. Equals to one if k genes changed and zero otherwise.
- void counter\_gains\_from\_0 (PhyloCounters \*counters, std::vector < size\_t > nfun, size\_t duplication=Geese::etype\_default)

  Used when all the functions are in 0 (like the root node prob.)
- void counter\_overall\_gains\_from\_0 (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

  Used when all the functions are in 0 (like the root node prob.)
- void counter\_pairwise\_overall\_change (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

  Used when all the functions are in 0 (like the root node prob.)
- void counter\_pairwise\_preserving (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_ 
  t duplication=Geese::etype\_default)

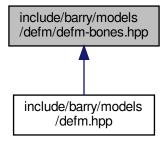
Used when all the functions are in 0 (like the root node prob.)

• void counter\_pairwise\_first\_gain (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_ t duplication=Geese::etype\_default)

Used when all the functions are in 0 (like the root node prob.)

# 9.35 include/barry/models/defm/defm-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:

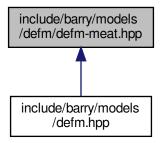


### Classes

class DEFM

# 9.36 include/barry/models/defm/defm-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



### **Macros**

- #define DEFM\_RANGES(a)
- #define DEFM\_LOOP\_ARRAYS(a) for (size\_t a = 0u; a < (nobs\_i M\_order); ++a)</li>

### **Functions**

• std::vector< double > keygen\_defm (const DEFMArray &Array\_, DEFMCounterData \*data)

### 9.36.1 Macro Definition Documentation

# 9.36.1.1 DEFM\_LOOP\_ARRAYS

Definition at line 36 of file defm-meat.hpp.

# 9.36.1.2 DEFM\_RANGES

Definition at line 31 of file defm-meat.hpp.

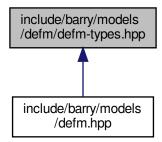
#### 9.36.2 Function Documentation

### 9.36.2.1 keygen\_defm()

Definition at line 4 of file defm-meat.hpp.

# 9.37 include/barry/models/defm/defm-types.hpp File Reference

This graph shows which files directly or indirectly include this file:



### **Classes**

- · class DEFMData
  - Data class for DEFM arrays.
- class DEFMCounterData

Data class used to store arbitrary size\_t or double vectors.

- class DEFMRuleData
- · class DEFMRuleDynData

# **Typedefs**

typedef barry::BArrayDense< int, DEFMData > DEFMArray

### Convenient typedefs for network objects.

- typedef barry::Counter< DEFMArray, DEFMCounterData > DEFMCounter
- typedef barry::Counters < DEFMArray, DEFMCounterData > DEFMCounters
- typedef barry::Support
   DEFMArray, DEFMCounterData, DEFMRuleData, DEFMRuleDynData > DEFMSupport
- typedef barry::StatsCounter< DEFMArray, DEFMCounterData > DEFMStatsCounter
- typedef barry::Model DEFMArray, DEFMCounterData, DEFMRuleData, DEFMRuleDynData > DEFMModel
- $\bullet \ \ type def \ barry:: Rule < DEFMArray, \ DEFMRule Data > DEFMRule \\$
- typedef barry::Rules < DEFMArray, DEFMRuleData > DEFMRules
- typedef barry::Rule < DEFMArray, DEFMRuleDynData > DEFMRuleDyn
- typedef barry::Rules < DEFMArray, DEFMRuleDynData > DEFMRulesDyn

# 9.37.1 Typedef Documentation

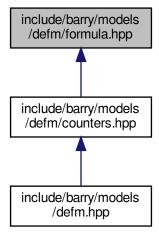
### 9.37.1.1 **DEFMArray**

typedef barry::BArrayDense<int, DEFMData> DEFMArray

Definition at line 5 of file defm-types.hpp.

# 9.38 include/barry/models/defm/formula.hpp File Reference

This graph shows which files directly or indirectly include this file:



### **Functions**

void defm\_motif\_parser (std::string formula, std::vector< size\_t > &locations, std::vector< bool > &signs, size\_t m\_order, size\_t y\_ncol)

Parses a motif formula.

### 9.38.1 Function Documentation

### 9.38.1.1 defm\_motif\_parser()

Parses a motif formula.

This function will take the formula and generate the corresponding input for defm::counter\_transition(). Formulas can be specified in the following ways:

- Intercept effect: {...} No transition, only including the current state.
- Transition effect: {...} > {...} Includes current and previous states.

The general notation is  $[0]y[column id]_[row id]$ . A preceeding zero means that the value of the cell is considered to be zero. The column id goes between 0 and the number of columns in the array - 1 (so it is indexed from 0,) and the row id goes from 0 to m\_order.

### Intercept effects

Intercept effects only involve a single set of curly brackets. Using the 'greater-than' symbol (i.e., '<') is only for transition effects. When specifying intercept effects, users can skip the  $row_id$ , e.g.,  $y0_0$  is equivalent to y0. If the passed  $row_id$  is different from the Markov order, i.e.,  $row_id$  !=  $m_order$ , then the function returns with an error.

### Examples:

• " $\{y0, 0y1\}$ " is equivalent to set a motif with the first element equal to one and the second to zero.

### **Transition effects**

Transition effects can be specified using two sets of curly brackets and an greater-than symbol, i.e.,  $\{\ldots\}$  >  $\{\ldots\}$ . The first set of brackets, which we call LHS, can only hold row id that are less than m\_order.

### **Parameters**

Definition at line 46 of file formula.hpp.

# 9.39 include/barry/models/geese.hpp File Reference

```
#include "geese/geese-types.hpp"
#include "geese/geese-node-bones.hpp"
#include "geese/geese-bones.hpp"
#include "geese/geese-meat.hpp"
#include "geese/geese-meat-constructors.hpp"
#include "geese/geese-meat-likelihood.hpp"
#include "geese/geese-meat-likelihood_exhaust.hpp"
#include "geese/geese-meat-simulate.hpp"
#include "geese/geese-meat-predict.hpp"
#include "geese/geese-meat-predict_exhaust.hpp"
#include "geese/geese-meat-predict_sim.hpp"
#include "geese/flock-bones.hpp"
#include "geese/flock-meat.hpp"
#include "geese/counters.hpp"
#include dependency graph for geese.hpp:
```



# **Namespaces**

• geese

# 9.40 include/barry/models/geese/flock-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



# **Classes**

· class Flock

A Flock is a group of Geese.

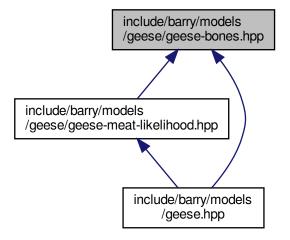
# 9.41 include/barry/models/geese/flock-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



# 9.42 include/barry/models/geese/geese-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



### Classes

· class Geese

Annotated Phylo Model.

### **Macros**

• #define INITIALIZED()

### **Functions**

```
    template < typename Ta , typename Tb > std::vector < Ta > vector_caster (const std::vector < Tb > &x)
    RULE_FUNCTION (rule_empty_free)
    std::vector < double > keygen_full (const PhyloArray & array, const PhyloCounterData *d)
```

bool vec diff (const std::vector< size t > &s, const std::vector< size t > &a)

# 9.42.1 Macro Definition Documentation

### **9.42.1.1 INITIALIZED**

```
#define INITIALIZED( )

Value:
    if (!this->initialized) \
    throw std::logic_error("The model has not been initialized yet.");
```

Definition at line 22 of file geese-bones.hpp.

### 9.42.2 Function Documentation

# 9.42.2.1 keygen\_full()

Definition at line 36 of file geese-bones.hpp.

### 9.42.2.2 RULE\_FUNCTION()

Definition at line 26 of file geese-bones.hpp.

# 9.42.2.3 vec\_diff()

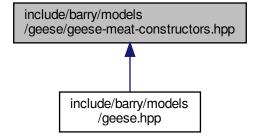
Definition at line 71 of file geese-bones.hpp.

# 9.42.2.4 vector\_caster()

Definition at line 10 of file geese-bones.hpp.

# 9.43 include/barry/models/geese/geese-meat-constructors.hpp File Reference

This graph shows which files directly or indirectly include this file:

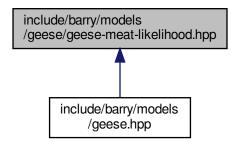


# 9.44 include/barry/models/geese/geese-meat-likelihood.hpp File Reference

#include "geese-bones.hpp"
Include dependency graph for geese-meat-likelihood.hpp:



This graph shows which files directly or indirectly include this file:



### **Functions**

void pset\_loop (size\_t n, size\_t s, size\_t nfunctions, const size\_t node\_id, const size\_t array\_id, std::vector< double > &totprob\_n, const std::vector< double > &par0, const std::vector< std::vector< bool >> &states, const std::vector< PhyloArray > &psets, const std::vector< std::vector< size\_t >> &locations, const std::vector< geese::Node \* > &node\_offspring, const double \*psetprobs)

# 9.44.1 Function Documentation

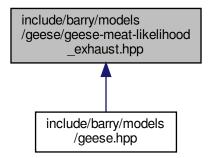
### 9.44.1.1 pset\_loop()

```
void pset_loop (
    size_t n,
    size_t s,
    size_t nfunctions,
    const size_t node_id,
    const size_t array_id,
    std::vector< double > & totprob_n,
    const std::vector< double > & par0,
    const std::vector< std::vector< bool >> & states,
    const std::vector< PhyloArray > & psets,
    const std::vector< std::vector< size_t > > & locations,
    const std::vector< geese::Node * > & node_offspring,
    const double * psetprobs ) [inline]
```

Definition at line 6 of file geese-meat-likelihood.hpp.

# 9.45 include/barry/models/geese/geese-meat-likelihood\_exhaust.hpp File Reference

This graph shows which files directly or indirectly include this file:



# 9.46 include/barry/models/geese/geese-meat-predict.hpp File Reference

This graph shows which files directly or indirectly include this file:



# 9.47 include/barry/models/geese/geese-meat-predict\_exhaust.hpp File Reference

This graph shows which files directly or indirectly include this file:



# 9.48 include/barry/models/geese/geese-meat-predict\_sim.hpp File Reference

This graph shows which files directly or indirectly include this file:



# 9.49 include/barry/models/geese/geese-meat-simulate.hpp File Reference

This graph shows which files directly or indirectly include this file:



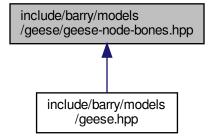
# 9.50 include/barry/models/geese/geese-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



# 9.51 include/barry/models/geese/geese-node-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



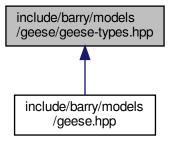
#### Classes

• class Node

A single node for the model.

# 9.52 include/barry/models/geese/geese-types.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Classes**

- · class NodeData
  - Data definition for the PhyloArray class.
- class PhyloCounterData
- · class PhyloRuleDynData

#### **Macros**

• #define POS(a, b) (b)\*N + (a)

## **Typedefs**

#### Convenient typedefs for Node objects.

- typedef std::vector< std::pair< size t, size t >> PhyloRuleData
- typedef barry::BArrayDense< size\_t, NodeData > PhyloArray
- typedef barry::Counter< PhyloArray, PhyloCounterData > PhyloCounter
- typedef barry::Counters
   PhyloArray, PhyloCounterData > PhyloCounters
- typedef barry::Rule < PhyloArray, PhyloRuleData > PhyloRule
- typedef barry::Rules
   PhyloArray, PhyloRuleData
   PhyloRules
- typedef barry::Rule < PhyloArray, PhyloRuleDynData > PhyloRuleDyn
- typedef barry::Rules
   PhyloArray, PhyloRuleDynData
   PhyloRulesDyn
- typedef barry::Support
   PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloSupport
- typedef barry::StatsCounter< PhyloArray, PhyloCounterData > PhyloStatsCounter
- typedef barry::Model < PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloModel
- typedef barry::PowerSet
   PhyloArray, PhyloRuleData > PhyloPowerSet

#### 9.52.1 Macro Definition Documentation

#### 9.52.1.1 POS

```
#define POS(  a, \\ b ) (b)*N + (a)
```

Definition at line 4 of file geese-types.hpp.

## 9.52.2 Typedef Documentation

#### 9.52.2.1 PhyloArray

```
typedef barry::BArrayDense<size_t, NodeData> PhyloArray
```

Definition at line 103 of file geese-types.hpp.

#### 9.52.2.2 PhyloCounter

```
typedef barry::Counter<PhyloArray, PhyloCounterData > PhyloCounter
```

Definition at line 104 of file geese-types.hpp.

#### 9.52.2.3 PhyloCounters

```
{\tt typedef\ barry::Counters} < {\tt PhyloArray,\ PhyloCounterData} > {\tt PhyloCounters}
```

Definition at line 105 of file geese-types.hpp.

#### 9.52.2.4 PhyloModel

 ${\tt typedef\ barry::Model < PhyloArray,\ PhyloCounterData,\ PhyloRuleData,\ PhyloRuleDynData > PhyloModel}$ 

Definition at line 115 of file geese-types.hpp.

#### 9.52.2.5 PhyloPowerSet

typedef barry::PowerSet<PhyloArray, PhyloRuleData> PhyloPowerSet

Definition at line 116 of file geese-types.hpp.

#### 9.52.2.6 PhyloRule

typedef barry::Rule<PhyloArray,PhyloRuleData> PhyloRule

Definition at line 107 of file geese-types.hpp.

#### 9.52.2.7 PhyloRuleData

typedef std::vector< std::pair< size\_t, size\_t > > PhyloRuleData

Definition at line 100 of file geese-types.hpp.

## 9.52.2.8 PhyloRuleDyn

typedef barry::Rule<PhyloArray,PhyloRuleDynData> PhyloRuleDyn

Definition at line 110 of file geese-types.hpp.

## 9.52.2.9 PhyloRules

typedef barry::Rules<PhyloArray,PhyloRuleData> PhyloRules

Definition at line 108 of file geese-types.hpp.

#### 9.52.2.10 PhyloRulesDyn

typedef barry::Rules<PhyloArray,PhyloRuleDynData> PhyloRulesDyn

Definition at line 111 of file geese-types.hpp.

#### 9.52.2.11 PhyloStatsCounter

typedef barry::StatsCounter<PhyloArray, PhyloCounterData> PhyloStatsCounter

Definition at line 114 of file geese-types.hpp.

#### 9.52.2.12 PhyloSupport

typedef barry::Support<PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloSupport

Definition at line 113 of file geese-types.hpp.

# 9.53 include/barry/powerset-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Classes**

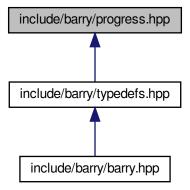
# 9.54 include/barry/powerset-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



# 9.55 include/barry/progress.hpp File Reference

This graph shows which files directly or indirectly include this file:



## **Classes**

• class Progress

A simple progress bar.

#### **Macros**

• #define BARRY\_PROGRESS\_BAR\_WIDTH 80

#### 9.55.1 Macro Definition Documentation

#### 9.55.1.1 BARRY PROGRESS BAR WIDTH

```
#define BARRY_PROGRESS_BAR_WIDTH 80
```

Definition at line 5 of file progress.hpp.

# 9.56 include/barry/rules-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



## **Classes**

- class Rule < Array\_Type, Data\_Type >
   Rule for determining if a cell should be included in a sequence.
- class Rules < Array\_Type, Data\_Type >

Vector of objects of class Rule.

#### **Functions**

template<typename Array\_Type , typename Data\_Type >
 bool rule\_fun\_default (const Array\_Type \*array, size\_t i, size\_t j, Data\_Type \*dat)

#### 9.56.1 Function Documentation

#### 9.56.1.1 rule\_fun\_default()

Definition at line 5 of file rules-bones.hpp.

# 9.57 include/barry/rules-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



# 9.58 include/barry/statscounter-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



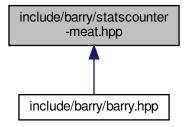
#### Classes

 $\bullet \ \ {\it class StatsCounter} < {\it Array\_Type}, \ {\it Data\_Type} >$ 

Count stats for a single Array.

# 9.59 include/barry/statscounter-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Macros**

- #define STATSCOUNTER TYPE() StatsCounter<Array Type, Data Type>
- #define STATSCOUNTER\_TEMPLATE\_ARGS() < typename Array\_Type, typename Data\_Type >
- #define STATSCOUNTER\_TEMPLATE(a, b) template STATSCOUNTER\_TEMPLATE\_ARGS() inline a STATSCOUNTER\_TYPE()::b

#### **Functions**

- STATSCOUNTER\_TEMPLATE (, StatsCounter)(const StatsCounter < Array\_Type
- EmptyArray clear ()
- STATSCOUNTER\_TEMPLATE (,~StatsCounter)()
- STATSCOUNTER\_TEMPLATE (void, reset\_array)(const Array\_Type \*Array\_)
- STATSCOUNTER\_TEMPLATE (void, add\_counter)(Counter< Array\_Type</li>
- STATSCOUNTER\_TEMPLATE (void, set\_counters)(Counters< Array\_Type
- STATSCOUNTER\_TEMPLATE (void, count\_init)(size\_t i
- current\_stats resize (counters->size(), 0.0)
- for (size\_t n=0u;n< counters->size();++n) current\_stats[n]
- STATSCOUNTER\_TEMPLATE (void, count\_current)(size\_t i
- STATSCOUNTER\_TEMPLATE (std::vector< std::string >, get\_names)() const
- STATSCOUNTER\_TEMPLATE (std::vector< std::string >, get\_descriptions)() const

#### **Variables**

```
    Data_Type & counter

EmptyArray = *Array
• current_stats = counter.current_stats

    counters = new Counters<Array_Type,Data_Type>((*counter.counters))
```

- counter\_deleted = false
- Data\_Type f\_
- return
- Data\_Type \* counters\_
- size\_t j

#### 9.59.1 Macro Definition Documentation

#### 9.59.1.1 STATSCOUNTER\_TEMPLATE

```
#define STATSCOUNTER_TEMPLATE(
             b) template STATSCOUNTER_TEMPLATE_ARGS() inline a STATSCOUNTER_TYPE()::b
```

Definition at line 8 of file statscounter-meat.hpp.

#### 9.59.1.2 STATSCOUNTER\_TEMPLATE\_ARGS

```
template STATSCOUNTER_TEMPLATE_ARGS( ) <typename Array_Type, typename Data_Type>
```

Definition at line 6 of file statscounter-meat.hpp.

#### 9.59.1.3 STATSCOUNTER TYPE

```
template Data_Type * STATSCOUNTER_TYPE( ) StatsCounter<Array_Type,Data_Type>
```

Definition at line 4 of file statscounter-meat.hpp.

#### 9.59.2 Function Documentation

#### 9.59.2.1 clear()

```
EmptyArray clear ( )
```

#### 9.59.2.2 for()

#### 9.59.2.3 resize()

## 9.59.2.4 STATSCOUNTER\_TEMPLATE() [1/9]

```
STATSCOUNTER_TEMPLATE (
StatsCounter ) const
```

#### 9.59.2.5 STATSCOUNTER\_TEMPLATE() [2/9]

```
STATSCOUNTER_TEMPLATE ( \sim \textit{StatsCounter} \ )
```

Definition at line 27 of file statscounter-meat.hpp.

## 9.59.2.6 STATSCOUNTER\_TEMPLATE() [3/9]

```
STATSCOUNTER_TEMPLATE (
          std::vector< std::string > ,
          get_descriptions ) const
```

Definition at line 256 of file statscounter-meat.hpp.

## 9.59.2.7 STATSCOUNTER\_TEMPLATE() [4/9]

```
STATSCOUNTER_TEMPLATE (
          std::vector< std::string > ,
          get_names ) const
```

Definition at line 251 of file statscounter-meat.hpp.

## 9.59.2.8 STATSCOUNTER\_TEMPLATE() [5/9]

## 9.59.2.9 STATSCOUNTER\_TEMPLATE() [6/9]

#### 9.59.2.10 STATSCOUNTER\_TEMPLATE() [7/9]

#### 9.59.2.11 STATSCOUNTER\_TEMPLATE() [8/9]

```
STATSCOUNTER_TEMPLATE (
void ,
reset_array ) const
```

Definition at line 34 of file statscounter-meat.hpp.

#### 9.59.2.12 STATSCOUNTER\_TEMPLATE() [9/9]

#### 9.59.3 Variable Documentation

#### 9.59.3.1 counter

```
Data_Type& counter

Initial value:
{
    Array = counter.Array
```

Definition at line 12 of file statscounter-meat.hpp.

## 9.59.3.2 counter\_deleted

```
counter_deleted = false
```

Definition at line 23 of file statscounter-meat.hpp.

#### 9.59.3.3 counters

```
counters = new Counters<Array_Type,Data_Type>((*counter.counters))
```

Definition at line 22 of file statscounter-meat.hpp.

#### 9.59.3.4 counters\_

```
Data_Type* counters_
Initial value:
{
    if (!counter_deleted)
        delete counters
```

Definition at line 53 of file statscounter-meat.hpp.

## 9.59.3.5 current\_stats

```
current_stats = counter.current_stats
```

Definition at line 19 of file statscounter-meat.hpp.

#### 9.59.3.6 EmptyArray

```
EmptyArray = *Array
```

Definition at line 17 of file statscounter-meat.hpp.

```
9.59.3.7 f
```

```
Data_Type f_
Initial value:
{
    counters->add_counter(f_)
```

Definition at line 44 of file statscounter-meat.hpp.

```
9.59.3.8 j
```

```
size_t j
Initial value:
{
    if (counters->size() == 0u)
```

throw std::logic\_error("No counters added: Cannot count without knowning what to count!")

Definition at line 66 of file statscounter-meat.hpp.

#### 9.59.3.9 return

return

Definition at line 49 of file statscounter-meat.hpp.

# 9.60 include/barry/support-bones.hpp File Reference

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.

# 9.61 include/barry/support-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



### **Macros**

• #define BARRY\_SUPPORT\_MEAT\_HPP 1

#### 9.61.1 Macro Definition Documentation

#### 9.61.1.1 BARRY\_SUPPORT\_MEAT\_HPP

#define BARRY\_SUPPORT\_MEAT\_HPP 1

Definition at line 2 of file support-meat.hpp.

# 9.62 include/barry/typedefs.hpp File Reference

```
#include "barry-configuration.hpp"
#include "barry-debug.hpp"
#include "progress.hpp"
Include dependency graph for typedefs.hpp:
```



This graph shows which files directly or indirectly include this file:



#### **Classes**

- class Entries < Cell\_Type >
  - A wrapper class to store source, target, val from a BArray object.
- struct vecHasher< 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 std::vector< std::pair< std::vector< double >, size t >> Counts type

    template<typename Cell_Type >
      using Row_type = Map< size_t, Cell< Cell_Type > >
    template<typename Cell_Type >
      using Col_type = Map< size_t, Cell< Cell_Type > * >
    • template<typename Ta = double, typename Tb = size_t>
      using MapVec_type = std::unordered_map< std::vector< Ta >, Tb, vecHasher< Ta >>
    • template<typename Array_Type , typename Data_Type >
      using Hasher_fun_type = std::function < std::vector < double >(const Array_Type &, Data_Type *)>
          Hasher function used by the counter.

    template<typename Array_Type , typename Data_Type >

      using Counter_fun_type = std::function < double(const Array_Type &, size_t, bata_Type &)>
          Counter and rule functions.
    • template<typename Array_Type , typename Data_Type >
      using Rule_fun_type = std::function< bool(const Array_Type &, size_t, Size_t, Data_Type &)>
Functions

    std::vector < size_t > sort_array (const double *v, size_t start, size_t ncols, size_t nrows)

          Ascending sorting an array.
    • template<typename T >
      T vec_inner_prod (const T *a, const T *b, size_t n)
    • template<> double vec_inner_prod (const double *a, const double *b, size_t n)
    template<typename T >
      bool vec equal (const std::vector< T > &a, const std::vector< T > &b)
          Compares if -a- and -b- are equal.
    • template<typename T >
      bool vec_equal_approx (const std::vector< T > &a, const std::vector< T > &b, double eps=1e-100)
    • const int CHECK::BOTH = -1
    • const int CHECK::NONE = 0
```

#### **Variables**

```
    const int CHECK::ONE = 1

    const int CHECK::TWO = 2

    const int EXISTS::BOTH = -1

    const int EXISTS::NONE = 0

• const int EXISTS::ONE = 1
• const int EXISTS::TWO = 1
• const int EXISTS::UKNOWN = -1
• const int EXISTS::AS ZERO = 0
const int EXISTS::AS_ONE = 1
```

## 9.62.1 Typedef Documentation

#### 9.62.1.1 Col type

```
template<typename Cell_Type >
using Col_type = Map< size_t, Cell<Cell_Type>* >
```

Definition at line 70 of file typedefs.hpp.

#### 9.62.1.2 Counter\_fun\_type

```
template<typename Array_Type , typename Data_Type >
using Counter_fun_type = std::function<double(const Array_Type &, size_t, size_t, Data_Type &)>
```

Counter and rule functions.

#### **Parameters**

Array_Type	a BArray
unit,size⇔	Focal cell
_t	
Data_Type	Data associated with the function, for example, id of the attribute in the Array.

#### Returns

```
Counter_fun_type a double (the change statistic) Rule_fun_type a bool. True if the cell is blocked.
```

Definition at line 187 of file typedefs.hpp.

#### 9.62.1.3 Counts\_type

```
typedef std::vector< std::pair< std::vector<double>, size_t >> Counts_type
Definition at line 51 of file typedefs.hpp.
```

#### 9.62.1.4 Hasher\_fun\_type

```
template<typename Array_Type , typename Data_Type >
using Hasher_fun_type = std::function<std::vector<double>(const Array_Type &, Data_Type *)>
```

Hasher function used by the counter.

Used to characterize the support of the array.

**Template Parameters** 

```
Array_Type
```

Definition at line 200 of file typedefs.hpp.

#### 9.62.1.5 MapVec\_type

```
template<typename Ta = double, typename Tb = size_t>
using MapVec_type = std::unordered_map< std::vector< Ta >, Tb, vecHasher<Ta> >
```

Definition at line 128 of file typedefs.hpp.

#### 9.62.1.6 Row\_type

```
template<typename Cell_Type >
using Row_type = Map< size_t, Cell<Cell_Type> >
```

Definition at line 67 of file typedefs.hpp.

#### 9.62.1.7 Rule\_fun\_type

```
template<typename Array_Type , typename Data_Type >
using Rule_fun_type = std::function<bool(const Array_Type &, size_t, size_t, Data_Type &)>
```

Definition at line 190 of file typedefs.hpp.

### 9.62.2 Function Documentation

#### 9.62.2.1 sort\_array()

Ascending sorting an array.

It will sort an array solving ties using the next column. Data is stored column-wise.

#### **Template Parameters**



#### **Parameters**



#### Returns

std::vector<size\_t> The sorting index.

Definition at line 141 of file typedefs.hpp.

#### 9.62.2.2 vec\_equal()

Compares if -a- and -b- are equal.

#### **Parameters**

```
a,b Two vectors of the same length
```

#### Returns

true if all elements are equal.

Definition at line 210 of file typedefs.hpp.

#### 9.62.2.3 vec\_equal\_approx()

Definition at line 235 of file typedefs.hpp.

## 9.62.2.4 vec\_inner\_prod() [1/2]

Definition at line 286 of file typedefs.hpp.

#### 9.62.2.5 vec\_inner\_prod() [2/2]

Definition at line 263 of file typedefs.hpp.

# 9.63 README.md File Reference

# Index

```
\simBArray
                                                           NetworkData, 205
    BArray< Cell Type, Data Type >, 63
                                                      \simNode
                                                           Node, 208
\simBArrayCell
    BArrayCell< Cell_Type, Data_Type >, 74
                                                      \simPhyloRuleDynData
~BArrayCell const
                                                           PhyloRuleDynData, 217
    BArrayCell_const< Cell_Type, Data_Type >, 77
                                                      \simPowerSet
{\sim}\mathsf{BArrayDense}
                                                           PowerSet < Array_Type, Data_Rule_Type >, 221
    BArrayDense < Cell_Type, Data_Type >, 83
                                                      \simProgress
\simBArrayDenseCell
                                                           Progress, 226
    BArrayDenseCell< Cell_Type, Data_Type >, 97
                                                      \simRule
\simBArrayRow
                                                           Rule < Array_Type, Data_Type >, 227
    BArrayRow< Cell Type, Data Type >, 110
                                                      \simRules
~BArrayRow const
                                                           Rules < Array Type, Data Type >, 230
    BArrayRow_const < Cell_Type, Data_Type >, 112
                                                      \simStatsCounter
\simBArrayVector
                                                           StatsCounter < Array_Type, Data_Type >, 235
    BArrayVector< Cell Type, Data Type >, 115
                                                      \simSupport
                                                                                       Data Counter Type,
~BArrayVector const
                                                           Support<
                                                                        Array_Type,
    BArrayVector_const< Cell_Type, Data_Type >,
                                                                Data_Rule_Type, Data_Rule_Dyn_Type >,
                                                                240
         118
\simCell
                                                      active
    Cell< Cell_Type >, 122
                                                           Cell< Cell_Type >, 125
\simConstBArrayRowIter
    ConstBArrayRowlter< Cell_Type, Data_Type >,
                                                      add
                                                           Cell < Cell_Type >, 123, 124
         127
                                                           FreqTable< T >, 158
\simCounter
                                                      add array
    Counter< Array_Type, Data_Type >, 130
                                                                                       Data Counter_Type,
                                                           Model <
                                                                       Array Type,
\simCounters
                                                                Data_Rule_Type, Data_Rule_Dyn_Type >,
    Counters < Array_Type, Data_Type >, 135
                                                                180
\simDEFMCounterData
                                                      add counter
    Phylo rules, 43
                                                           Counters < Array_Type, Data_Type >, 135, 136
\simDEFMData
                                                           Model <
                                                                       Array_Type,
                                                                                       Data_Counter_Type,
    Phylo rules, 43
                                                                Data_Rule_Type, Data_Rule_Dyn_Type >,
\simDEFMRuleDynData
    Phylo rules, 43
                                                           StatsCounter < Array_Type, Data_Type >, 235
\simEntries
                                                           Support<
                                                                        Array_Type,
                                                                                       Data_Counter_Type,
    Entries < Cell Type >, 149
                                                                Data_Rule_Type, Data_Rule_Dyn_Type >,
\simFlock
                                                                240
    Flock, 151
                                                      add data
\simFreqTable
                                                           Flock, 151
    FreqTable < T >, 158
                                                      add dims
\simGeese
                                                           counters-meat.hpp, 287
    Geese, 164
                                                      add hash
\simModel
                                                           Counters < Array Type, Data Type >, 136
    Model<
                Array Type,
                                 Data Counter Type,
         Data_Rule_Type, Data_Rule_Dyn_Type >,
                                                      add hasher
                                                           Model<
                                                                                       Data Counter Type,
                                                                       Array Type,
         180
                                                                Data_Rule_Type, Data_Rule_Dyn_Type >,
\simNetCounterData
    NetCounterData, 203
                                                      add_rule
\simNetworkData
```

Model< Array_Type, Data_Counter_Type,	ncol, 67
Data_Rule_Type, Data_Rule_Dyn_Type >,	nnozero, 67
181, 182	nrow, 67
PowerSet < Array_Type, Data_Rule_Type >, 221	operator*=, 67
Rules < Array_Type, Data_Type >, 231	operator(), 67
Support< Array_Type, Data_Counter_Type,	operator+=, 68
Data_Rule_Type, Data_Rule_Dyn_Type >,	operator-=, 68
240, 241	operator/=, 69
	operator=, 69
add_rule_dyn  Model< Array Type. Data Counter Type.	operator==, 69
	•
Data_Rule_Type, Data_Rule_Dyn_Type >,	out_of_range, 69
182	print, 69
Support< Array_Type, Data_Counter_Type,	print_n, 70
Data_Rule_Type, Data_Rule_Dyn_Type >,	reserve, 70
241	resize, 70
annotations	rm_cell, 70
Node, 209	row, 70
Array	set_data, 71
ConstBArrayRowIter< Cell_Type, Data_Type >,	swap_cells, 71
127	swap_cols, 71
array	swap_rows, 71
Node, 209	toggle_cell, 72
Phylo rules, 44	toggle_lock, 72
arrays	transpose, 72
Node, 209	visited, 73
arrays2support	zero_col, 72
Model< Array_Type, Data_Counter_Type,	zero_row, 72
Data_Rule_Type, Data_Rule_Dyn_Type >,	
195	BARRAY_TEMPLATE, 252-254
AS_ONE	BARRAY_TEMPLATE_ARGS, 253, 255
EXISTS, 57	BARRAY_TYPE, 253, 255
as vector	COL, 253
FreqTable< T >, 158	for, 255
AS_ZERO	operator(), 255
EXISTS, 57	rhs, 255
at	ROW, 253
Phylo rules, 39	this, 256
PhyloCounterData, 214	barray-meat.hpp
Thylocounterbata, 214	COL, 256
BArray	
BArray< Cell_Type, Data_Type >, 62, 63	ROW, 257
BArray< Cell_Type, Data_Type >, 59	BARRAY_TEMPLATE
~BArray, 63	barray-meat-operators.hpp, 252–254
BArray, 62, 63	BARRAY_TEMPLATE_ARGS
BArrayCell< Cell_Type, Data_Type >, 73	barray-meat-operators.hpp, 253, 255
BArrayCell_const< Cell_Type, Data_Type >, 73	BARRAY_TYPE
clear, 63	barray-meat-operators.hpp, 253, 255
col, 63	BArrayCell
D, 64	BArrayCell< Cell_Type, Data_Type >, 74
	BArrayCell< Cell_Type, Data_Type >, 74
D_ptr, 64	$\sim$ BArrayCell, 74
default_val, 64	BArray< Cell_Type, Data_Type >, 73
flush_data, 64	BArrayCell, 74
get_cell, 64	operator Cell_Type, 75
get_col_vec, 65	operator*=, 75
get_entries, 65	operator+=, 75
get_row_vec, 65	operator-=, 75
insert_cell, 66	operator/=, 75
is_dense, 66	operator=, 75
is_empty, 66	•

operator==, 76	row, 92
BArrayCell_const	rowsum, 92
BArrayCell_const< Cell_Type, Data_Type >, 77	set_data, 93
BArrayCell_const< Cell_Type, Data_Type >, 76	swap_cells, 93
~BArrayCell_const, 77	swap_cols, 93
BArray< Cell_Type, Data_Type >, 73	swap_rows, 93
BArrayCell_const, 77	toggle_cell, 94
operator Cell_Type, 77	toggle_lock, 94
operator!=, 77	transpose, 94
operator<, 77	visited, 96
operator<=, 78	zero_col, 94
operator>, 78	zero_row, 94
operator>=, 78	barraydense-meat-operators.hpp
operator==, 78	BDENSE_TEMPLATE, 259, 261
BArrayDense	BDENSE_TEMPLATE_ARGS, 259, 261
BArrayDense< Cell_Type, Data_Type >, 82, 83	BDENSE_TYPE, 260, 262
BArrayDense< Cell_Type, Data_Type >, 79	COL, 260
$\sim$ BArrayDense, 83	POS, 260
BArrayDense, 82, 83	POS_N, 260
BArrayDenseCell< Cell_Type, Data_Type >, 95,	ROW, 260
99	barraydense-meat.hpp
BArrayDenseCol< Cell_Type, Data_Type >, 95,	COL, 262
102	POS, 262
BArrayDenseCol_const< Cell_Type, Data_Type >,	POS_N, 263
95	ROW, 263
BArrayDenseRow< Cell_Type, Data_Type >, 95,	ZERO_CELL, 263
106	BArrayDenseCell
BArrayDenseRow_const< Cell_Type, Data_Type	BArrayDenseCell< Cell_Type, Data_Type >, 97
>, 95	BArrayDenseCell< Cell_Type, Data_Type >, 96
clear, 83	$\sim$ BArrayDenseCell, 97
col, 84	BArrayDense < Cell_Type, Data_Type >, 95, 99
colsum, 84	BArrayDenseCell, 97
D, 84	BArrayDenseCol< Cell_Type, Data_Type >, 99,
D_ptr, 85	102
default_val, 85	BArrayDenseCol_const< Cell_Type, Data_Type >,
get_cell, 85	99, 104
get_col_vec, 85, 86	BArrayDenseRow< Cell_Type, Data_Type >, 106
get_data, 86	BArrayDenseRow_const< Cell_Type, Data_Type
get_entries, 86	>, 109
get_row_vec, 86, 87	operator Cell_Type, 97
insert_cell, 87	operator*=, 97
is_dense, 87	operator+=, 98
is_empty, 88	operator-=, 98
ncol, 88	operator/=, 98
nnozero, 88	operator=, 98
nrow, 88	operator==, 99
operator∗=, 89	barraydensecell-bones.hpp
operator(), 88, 89	POS, 264
operator+=, 89	barraydensecell-meat.hpp
operator-=, 90	POS, 264
operator/=, 90	BArrayDenseCell_const< Cell_Type, Data_Type >, 100
operator=, 90, 91	BArrayDenseCol< Cell_Type, Data_Type >, 102
operator==, 91	BArrayDenseCol_const< Cell_Type, Data_Type >,
out_of_range, 91	104
print, 91	BArrayDenseRow< Cell_Type, Data_Type >, 106
reserve, 91	BArrayDenseRow_const< Cell_Type, Data_Type
resize, 92	>, 109
rm_cell, 92	BArrayDenseCol

BArrayDenseCol< Cell_Type, Data_Type >, 100	operator(), 108
BArrayDenseCol Cell_Type, Data_Type >, 100	size, 108
BArrayDense Cell_Type, Data_Type >, 95, 102	BArrayRow
BArrayDenseCell< Cell_Type, Data_Type >, 99, 102	BArrayRow< Cell_Type, Data_Type >, 110 BArrayRow< Cell_Type, Data_Type >, 109
BArrayDenseCell_const< Cell_Type, Data_Type	~BArrayRow, 110
>, 102	BArrayRow, 110
BArrayDenseCol, 100	operator BArrayRow< Cell_Type, Data_Type >,
begin, 101	110
end, 101	operator*=, 110
operator(), 101	operator+=, 110
size, 101	operator-=, 110
barraydensecol-bones.hpp	operator/=, 111
POS, 265	operator=, 111
POS_N, 265	operator==, 111
ZERO_CELL, 266	barrayrow-meat.hpp
BArrayDenseCol_const	BROW_TEMPLATE, 268, 269
${\sf BArrayDenseCol\_const} {< {\sf Cell\_Type}, {\sf Data\_Type} >},$	BROW_TEMPLATE_ARGS, 268
103	BROW_TYPE, 268
BArrayDenseCol_const< Cell_Type, Data_Type >, 102	BArrayRow_const
BArrayDense< Cell_Type, Data_Type >, 95	BArrayRow_const< Cell_Type, Data_Type >, 112
BArrayDenseCell< Cell_Type, Data_Type >, 99,	BArrayRow_const< Cell_Type, Data_Type >, 111
104	~BArrayRow_const, 112
BArrayDenseCell_const< Cell_Type, Data_Type	BArrayRow_const, 112
>, 104	operator BArrayRow_const< Cell_Type, Data_Type
BArrayDenseCol_const, 103	>, 112
begin, 103	operator!=, 112
end, 103	operator<, 112
operator(), 103	operator<=, 113
size, 104 BArrayDenseRow	operator>, 113
BArrayDenseRow< Cell_Type, Data_Type >, 105	operator>=, 113
BArrayDenseRow< Cell_Type, Data_Type >, 104	operator==, 113 BArrayVector
BArrayDense< Cell_Type, Data_Type >, 95, 106	BArrayVector< Cell_Type, Data_Type >, 114
BArrayDenseCell< Cell_Type, Data_Type >, 106	BArrayVector< Cell_Type, Data_Type >, 113
BArrayDenseCell_const< Cell_Type, Data_Type  BArrayDenseCell_const< Cell_Type, Data_Type	~BArrayVector, 115
>, 106	BArrayVector, 114
BArrayDenseRow, 105	begin, 115
begin, 105	end, 115
end, 105	is_col, 115
operator(), 106	is_row, 115
size, 106	operator std::vector< Cell_Type >, 116
barraydenserow-bones.hpp	operator*=, 116
POS, 267	operator+=, 116
POS_N, 267	operator-=, 116
ZERO_CELL, 267	operator/=, 116
BArrayDenseRow_const	operator=, 117
BArrayDenseRow_const< Cell_Type, Data_Type	operator==, 117
>, 107	size, 117
BArrayDenseRow_const< Cell_Type, Data_Type >,	BArrayVector_const
107	BArrayVector_const< Cell_Type, Data_Type >,
BArrayDense Cell_Type, Data_Type >, 95	118
BArrayDenseCell Const Cell Type, Data_Type >, 109	BArrayVector_const < Cell_Type, Data_Type >, 117
BArrayDenseCell_const< Cell_Type, Data_Type >, 109	~BArrayVector_const, 118 BArrayVector_const, 118
>, 109 BArrayDenseRow_const, 107	begin, 119
begin, 108	end, 119
end, 108	is_col, 119
oria, roo	13_001, 110

is_row, 119	support-meat.hpp, 341
operator std::vector< Cell_Type >, 119	BARRY_UNUSED
operator!=, 119	barry-macros.hpp, 274
operator<, 120	BARRY_VERSION
operator<=, 120	barry.hpp, 276
operator>, 120	BARRY_VERSION_MAYOR
operator>=, 120	barry.hpp, 276
operator==, 120	BARRY_VERSION_MINOR
size, 121	barry.hpp, 276
barry, 55	BARRY_ZERO
barry-configuration.hpp	barry-macros.hpp, 274
BARRY_CHECK_SUPPORT, 271	BARRY_ZERO_DENSE
BARRY_ISFINITE, 271	barry-macros.hpp, 274
BARRY_MAX_NUM_ELEMENTS, 271	BARRY_ZERO_NETWORK
BARRY_SAFE_EXP, 271	network.hpp, 302
Map, 272	BARRY_ZERO_NETWORK_DENSE
printf_barry, 271	network.hpp, 302
barry-debug.hpp	BDENSE_TEMPLATE
BARRY_DEBUG_LEVEL, 272	barraydense-meat-operators.hpp, 259, 261
barry-macros.hpp	BDENSE_TEMPLATE_ARGS
BARRY_NCORES_ARG, 273	barraydense-meat-operators.hpp, 259, 261
BARRY_ONE, 273	BDENSE_TYPE
BARRY_ONE_DENSE, 273	barraydense-meat-operators.hpp, 260, 262
BARRY_UNUSED, 274	begin
BARRY_ZERO, 274	BArrayDenseCol< Cell_Type, Data_Type >, 101
BARRY_ZERO_DENSE, 274	BArrayDenseCol_const< Cell_Type, Data_Type >,
barry.hpp	103
BARRY_HPP, 276	BArrayDenseRow< Cell_Type, Data_Type >, 105
BARRY_VERSION, 276	BArrayDenseRow_const< Cell_Type, Data_Type
BARRY_VERSION_MAYOR, 276	>, 108
BARRY_VERSION_MINOR, 276	BArrayVector< Cell_Type, Data_Type >, 115
COUNTER_FUNCTION, 276	BArrayVector_const< Cell_Type, Data_Type >,
COUNTER_LAMBDA, 276	119
RULE_FUNCTION, 277	PhyloCounterData, 215
RULE_LAMBDA, 277	PowerSet< Array_Type, Data_Rule_Type >, 221
barry::counters, 55	Rules < Array_Type, Data_Type >, 231
barry::counters::network, 56	blengths
BARRY_CHECK_SUPPORT	NodeData, 212
barry-configuration.hpp, 271	ВОТН
BARRY_DEBUG_LEVEL	CHECK, 56
barry-debug.hpp, 272	EXISTS, 57
BARRY_HPP	BROW_TEMPLATE
barry.hpp, 276	barrayrow-meat.hpp, 268, 269
BARRY_ISFINITE	BROW_TEMPLATE_ARGS
barry-configuration.hpp, 271	barrayrow-meat.hpp, 268
BARRY_MAX_NUM_ELEMENTS	BROW_TYPE
barry-configuration.hpp, 271	barrayrow-meat.hpp, 268
BARRY_NCORES_ARG	calc
barry-macros.hpp, 273	PowerSet < Array_Type, Data_Rule_Type >, 221
BARRY_ONE	Support< Array_Type, Data_Counter_Type,  Support< Array_Type, Data_Counter_Type,
barry-macros.hpp, 273	
BARRY_ONE_DENSE	Data_Rule_Type, Data_Rule_Dyn_Type >, 241
barry-macros.hpp, 273	
BARRY_PROGRESS_BAR_WIDTH	calc_reduced_sequence Geese, 164
progress.hpp, 333	calc_sequence
BARRY_SAFE_EXP	Geese, 165
barry-configuration.hpp, 271	Cell
BARRY_SUPPORT_MEAT_HPP	Cell < Cell_Type >, 122, 123
	Oui < Oui_ Type /, 122, 120

Cell< Cell_Type >, 121	coordiantes_n_free
∼Cell, 122	Support< Array_Type, Data_Counter_Type
active, 125	Data_Rule_Type, Data_Rule_Dyn_Type > 245
add, 123, 124	
Cell, 122, 123	coordiantes_n_locked Support< Array Type, Data Counter Type
operator Cell_Type, 124	
operator!=, 124	Data_Rule_Type, Data_Rule_Dyn_Type >
operator=, 124, 125	246
operator==, 125	coordinates_free
value, 125	PowerSet < Array_Type, Data_Rule_Type >, 223
visited, 125	Support< Array_Type, Data_Counter_Type
Cell_const< Cell_Type >, 126	Data_Rule_Type, Data_Rule_Dyn_Type >
change_stats	246
Support< Array_Type, Data_Counter_Type,	
Data_Rule_Type, Data_Rule_Dyn_Type >,	PowerSet< Array_Type, Data_Rule_Type >, 223
245	Support< Array_Type, Data_Counter_Type
CHECK, 56	Data_Rule_Type, Data_Rule_Dyn_Type >
BOTH, 56	246
NONE, 56	count
ONE, 56	Counter< Array_Type, Data_Type >, 131
TWO, 56	count_all
clear	StatsCounter < Array_Type, Data_Type >, 235
BArray< Cell_Type, Data_Type >, 63	count_current
BArrayDense< Cell_Type, Data_Type >, 83	StatsCounter < Array_Type, Data_Type >, 236
FreqTable $<$ T $>$ , 158	count_fun
statscounter-meat.hpp, 336	Counter< Array_Type, Data_Type >, 132
COL	counters-meat.hpp, 282
barray-meat-operators.hpp, 253	count_fun_
barray-meat.hpp, 256	counters-meat.hpp, 287
barraydense-meat-operators.hpp, 260	count_init
barraydense-meat.hpp, 262	StatsCounter< Array_Type, Data_Type >, 236
col	Counter
BArray< Cell_Type, Data_Type >, 63	Counter< Array_Type, Data_Type >, 130
BArrayDense< Cell_Type, Data_Type >, 84	counter
Col_type	counters-meat.hpp, 288
typedefs.hpp, 344	statscounter-meat.hpp, 338
colnames	Counter< Array_Type, Data_Type >, 128
Flock, 152	$\sim$ Counter, 130
Geese, 165	count, 131
Model< Array_Type, Data_Counter_Type,	count_fun, 132
Data_Rule_Type, Data_Rule_Dyn_Type >,	Counter, 130
182	data, 132
colsum	desc, 133
BArrayDense< Cell_Type, Data_Type >, 84	get_description, 131
conditional_prob	get_hasher, 131
Model< Array_Type, Data_Counter_Type,	get_name, 131
Data_Rule_Type, Data_Rule_Dyn_Type >,	hasher_fun, 133
182	init, 131
ConstBArrayRowlter	init_fun, 133
ConstBArrayRowIter< Cell_Type, Data_Type >,	name, 133
127	operator=, 131, 132
ConstBArrayRowlter< Cell_Type, Data_Type >, 126	set_hasher, 132
~ConstBArrayRowlter, 127	counter_
Array, 127	counters-meat.hpp, 288
ConstBArrayRowlter, 127	counter_absdiff
current_col, 128	Network counters, 26
current_row, 128	counter_co_opt
iter, 128	Counting, 18

Dhyle counters 40	Dhula aquatara 40
Phylo counters, 48	Phylo counters, 49
counter_cogain	counter_gains_from_0
Counting, 18	Counting, 18
Phylo counters, 49	Phylo counters, 49
counter_css_census01	counter_gains_k_offspring
network-css.hpp, 295	Counting, 19
counter_css_census02	Phylo counters, 49
network-css.hpp, 295	counter_genes_changing
counter_css_census03	Counting, 19
network-css.hpp, 295	Phylo counters, 50
counter_css_census04	counter_idegree
network-css.hpp, 296	Network counters, 29
counter_css_census05	counter_idegree15
network-css.hpp, 296	Network counters, 29
counter_css_census06	counter_isolates
network-css.hpp, 296	Network counters, 29, 30
counter_css_census07	counter_istar2
network-css.hpp, 296	Network counters, 30
counter_css_census08	counter_k_genes_changing
network-css.hpp, 297	Counting, 19
counter_css_census09	Phylo counters, 50
network-css.hpp, 297	COUNTER_LAMBDA
counter_css_census10	barry.hpp, 276
network-css.hpp, 297	counter_less_than_p_prop_genes_changing
counter_css_completely_false_recip_comiss	Counting, 19
network-css.hpp, 297	Phylo counters, 50
counter_css_completely_false_recip_omiss	counter_logit_intercept
network-css.hpp, 298	Network counters, 30
counter_css_mixed_recip	counter_longest
network-css.hpp, 298	Counting, 20
counter_css_partially_false_recip_commi	Phylo counters, 50
network-css.hpp, 298	counter_loss
counter_css_partially_false_recip_omiss	Counting, 20
network-css.hpp, 299	Phylo counters, 51
counter_ctriads	counter_maxfuns
Network counters, 27	Counting, 20
counter_degree	Phylo counters, 51
Network counters, 27	counter_mutual
counter_deleted	Network counters, 30
statscounter-meat.hpp, 339	counter_neofun
counter_density	Counting, 20
Network counters, 27	Phylo counters, 51
counter_diff	counter_neofun_a2b
Network counters, 28	Counting, 21
counter_edges	Phylo counters, 51
Network counters, 28	counter_nodecov
counter_fixed_effect	Network counters, 31
Network counters, 28	counter_nodeicov
counter_fun	Network counters, 31
Model< Array_Type, Data_Counter_Type,	counter_nodematch
Data_Rule_Type, Data_Rule_Dyn_Type >,	Network counters, 31
195	counter_nodeocov
Counter_fun_type	Network counters, 31
typedefs.hpp, 344	counter_odegree
COUNTER_FUNCTION	Network counters, 31, 32
barry.hpp, 276	counter_odegree15
counter_gains	Network counters, 32
Counting, 18	counter ones

Network counters, 32	add_hash, 136
counter_ostar2	Counters, 134, 135
Network counters, 33	gen_hash, 136
counter_overall_changes	get_descriptions, 136
Counting, 21	get_names, 137
Phylo counters, 52	operator=, 137
counter_overall_gains	operator[], 138
Counting, 21	size, 138
Phylo counters, 52	counters-meat.hpp
counter_overall_gains_from_0	add_dims, 287
Counting, 21	count_fun, 282
Phylo counters, 52	count_fun_, 287
counter_overall_loss	counter, 288
Counting, 22	counter_, 288
Phylo counters, 52	COUNTER_TEMPLATE, 280, 282, 283
counter_pairwise_first_gain	COUNTER TEMPLATE ARGS, 280
Counting, 22	COUNTER TYPE, 281
Phylo counters, 53	COUNTERS TEMPLATE, 281, 283–285
counter_pairwise_neofun_singlefun	COUNTERS TEMPLATE ARGS, 281
Counting, 22	COUNTERS TYPE, 281
Phylo counters, 53	data, 285
counter_pairwise_overall_change	data_, 288
Counting, 22	desc, 285
Phylo counters, 53	desc , 288
counter_pairwise_preserving	for, 285
Counting, 23	fun, 289
Phylo counters, 53	fun_, 289
-	hasher, 285, 286
counter_preserve_pseudogene	
Counting, 23	hasher_fun, 286
Phylo counters, 54	hasher_fun_, 289
counter_prop_genes_changing	i, 289
Counting, 23	if, 286
Phylo counters, 54	init_fun, 287
counter_subfun	init_fun_, 290
Counting, 23	j, 290
Phylo counters, 54	name, 287
COUNTER_TEMPLATE	name_, 290
counters-meat.hpp, 280, 282, 283	noexcept, 290
COUNTER_TEMPLATE_ARGS	res, 290
counters-meat.hpp, 280	return, 291
counter_transition	TMP_HASHER_CALL, 281
Network counters, 33	counters.hpp
counter_transition_formula	DEFM_COUNTER, 310
Network counters, 34	DEFM_COUNTER_LAMBDA, 311
counter_ttriads	DEFM_RULE, 311
Network counters, 34	DEFM_RULE_LAMBDA, 311
COUNTER_TYPE	DEFM_RULEDYN_LAMBDA, 311
counters-meat.hpp, 281	UNI_SUB, 312
Counters	counters_
Counters < Array_Type, Data_Type >, 134, 135	statscounter-meat.hpp, 339
counters	COUNTERS_TEMPLATE
Model< Array_Type, Data_Counter_Type,	counters-meat.hpp, 281, 283–285
Data_Rule_Type, Data_Rule_Dyn_Type >,	COUNTERS_TEMPLATE_ARGS
195	counters-meat.hpp, 281
statscounter-meat.hpp, 339	COUNTERS_TYPE
Counters < Array_Type, Data_Type >, 134	counters-meat.hpp, 281
∼Counters, 135	Counting, 13
add counter, 135, 136	counter co opt. 18

counter_cogain, 18	CSS_MATCH_TYPE
counter_gains, 18	network-css.hpp, 294
counter_gains_from_0, 18	CSS_NET_COUNTER_LAMBDA_INIT
counter_gains_k_offspring, 19	network-css.hpp, 294
counter_genes_changing, 19	CSS_PERCEIVED_CELLS
counter_k_genes_changing, 19	network-css.hpp, 294
counter_less_than_p_prop_genes_changing, 19	CSS_SIZE
counter_longest, 20	network-css.hpp, 294
counter_loss, 20	CSS_TRUE_CELLS
counter_maxfuns, 20	network-css.hpp, 295
counter_neofun, 20	current_col
counter_neofun_a2b, 21	ConstBArrayRowlter< Cell Type, Data Type >,
counter_overall_changes, 21	128
counter_overall_gains, 21	current_row
counter_overall_gains_from_0, 21	ConstBArrayRowlter< Cell_Type, Data_Type >,
counter_overall_loss, 22	128
counter_pairwise_first_gain, 22	current stats
counter_pairwise_neofun_singlefun, 22	statscounter-meat.hpp, 339
counter_pairwise_overall_change, 22	Support< Array_Type, Data_Counter_Type,
counter pairwise preserving, 23	Data Rule Type, Data Rule Dyn Type >,
counter preserve pseudogene, 23	246
counter_prop_genes_changing, 23	240
counter_subfun, 23	D
get last name, 24	BArray< Cell_Type, Data_Type >, 64
IF_MATCHES, 15	BArrayDense< Cell_Type, Data_Type >, 84
IF_NOTMATCHES, 15	Rule < Array_Type, Data_Type >, 228
IS DUPLICATION, 15	D_ptr
IS EITHER, 15	BArray< Cell_Type, Data_Type >, 64
IS SPECIATION, 16	BArrayDense < Cell_Type, Data_Type >, 85
MAKE_DEFM_HASHER, 16	dat
MAKE_DUPL_VARS, 16	Flock, 156
PHYLO_CHECK_MISSING, 16	data
PHYLO_COUNTER_LAMBDA, 17	Counter< Array_Type, Data_Type >, 132
PHYLO_RULE_DYN_LAMBDA, 17	counters-meat.hpp, 285
PHYLO_RULE_LAMBDA, 17	PowerSet < Array_Type, Data_Rule_Type >, 223
counts	data_
Phylo rules, 44	counters-meat.hpp, 288
PhyloRuleDynData, 218	default_val
Counts_type	BArray< Cell_Type, Data_Type >, 64
typedefs.hpp, 344	BArrayDense < Cell_Type, Data_Type >, 85
covar sort	DEFM, 139
Phylo rules, 44	DEFM, 140
covar used	get_column_major, 140
Phylo rules, 44	get_ID, 140
covariates	get_m_order, 140
Phylo rules, 44	get_model, 141
CSS APPEND	get_n_covars, 141
network-css.hpp, 292	get_n_obs, 141
CSS_CASE_ELSE	get_n_rows, 141
network-css.hpp, 293	get_n_y, 141
CSS_CASE_PERCEIVED	get_X, 141
network-css.hpp, 293	get_X_names, 142
CSS_CASE_TRUTH	get_Y, 142
network-css.hpp, 293	get_Y_names, 142
CSS_CHECK_SIZE	init, 142
network-css.hpp, 293	is_motif, 142
CSS_CHECK_SIZE_INIT	logodds, 142
network-css.hpp, 293	motif_census, 143
117 - 77	print, 143

set_names, 143	Data Rule Type, Data Rule Dyn Type >,
simulate, 143	246
defm, 57	delete_rengine
defm-meat.hpp	Geese, 172
DEFM LOOP ARRAYS, 315	Model< Array_Type, Data_Counter_Type,
DEFM RANGES, 315	Data_Rule_Type, Data_Rule_Dyn_Type >,
keygen_defm, 316	195
defm-types.hpp	delete_rules
DEFMArray, 317	Model< Array_Type, Data_Counter_Type,
DEFM COUNTER	Data_Rule_Type, Data_Rule_Dyn_Type >,
counters.hpp, 310	196
DEFM_COUNTER_LAMBDA	Support< Array_Type, Data_Counter_Type,
counters.hpp, 311	Data_Rule_Type, Data_Rule_Dyn_Type >,
DEFM_LOOP_ARRAYS	247
defm-meat.hpp, 315	delete_rules_dyn
defm_motif_parser	Model < Array_Type, Data_Counter_Type,
formula.hpp, 317	Data_Rule_Type, Data_Rule_Dyn_Type >,
DEFM_RANGES	196
defm-meat.hpp, 315	Support< Array_Type, Data_Counter_Type,
DEFM_RULE	Data_Rule_Type, Data_Rule_Dyn_Type >, 247
counters.hpp, 311	
DEFM_RULE_LAMBDA	delete_support
counters.hpp, 311	Geese, 172
DEFM_RULEDYN_LAMBDA	desc
counters.hpp, 311	Counter < Array_Type, Data_Type >, 133
DEFMArray	counters-meat.hpp, 285
defm-types.hpp, 317	desc_
DEFMCounter	counters-meat.hpp, 288
Phylo rules, 37	directed
DEFMCounterData, 144	NetworkData, 205
Phylo rules, 39	duplication
DEFMCounters	Node, 210
Phylo rules, 37	NodeData, 213
DEFMData, 144	PhyloRuleDynData, 218
Phylo rules, 39	amatu
DEFMModel	empty  Phyla Countary Data 245
Phylo rules, 37	PhyloCounterData, 215
DEFMRule	EmptyArray
Phylo rules, 38	PowerSet < Array_Type, Data_Rule_Type >, 224
DEFMRuleData, 146	statscounter-meat.hpp, 339
Phylo rules, 40	end
DEFMRuleDyn	BArrayDenseCol< Cell_Type, Data_Type >, 101
Phylo rules, 38	BArrayDenseCol_const< Cell_Type, Data_Type >,
DEFMRuleDynData, 147	103
Phylo rules, 40	BArrayDenseRow< Cell_Type, Data_Type >, 105
DEFMRules	BArrayDenseRow_const< Cell_Type, Data_Type
Phylo rules, 38	>, 108
DEFMRulesDyn	BArrayVector< Cell_Type, Data_Type >, 115
Phylo rules, 38	$BArrayVector\_const < Cell\_Type, \;\; Data\_Type \;\; >,$
DEFMStatsCounter	119
Phylo rules, 38	PhyloCounterData, 215
DEFMSupport	PowerSet < Array_Type, Data_Rule_Type >, 222
Phylo rules, 38	Progress, 226
delete_counters	Rules < Array_Type, Data_Type >, 231
Model< Array_Type, Data_Counter_Type,	Entries
Data_Rule_Type, Data_Rule_Dyn_Type >,	Entries < Cell_Type >, 148
195	Entries < Cell_Type >, 148
Support< Array_Type, Data_Counter_Type,	$\sim$ Entries, 149
	Entrice 1/18

resize, 149	BArray< Cell_Type, Data_Type >, 64
source, 149	for
target, 149	barray-meat-operators.hpp, 255
val, 149	counters-meat.hpp, 285
etype_default	statscounter-meat.hpp, 336
Geese, 172	formula.hpp
etype_duplication	defm_motif_parser, 317
Geese, 172	FreqTable
etype_either	FreqTable< T >, 158
Geese, 172	FreqTable < T >, 157
etype_speciation	∼FregTable, 158
Geese, 173	add, 158
eval_rules_dyn	as_vector, 158
Support< Array_Type, Data_Counter_Type,	clear, 158
Data_Rule_Type, Data_Rule_Dyn_Type >,	FreqTable, 158
242	get_data, 159
EXISTS, 57	get_index, 159
AS ONE, 57	make_hash, 159
AS ZERO, 57	print, 159
BOTH, 57	reserve, 159
NONE, 58	size, 160
ONE, 58	fun
TWO, 58	counters-meat.hpp, 289
UKNOWN, 58	fun_
UNIVOVIV, 56	
f_	counters-meat.hpp, 289
statscounter-meat.hpp, 340	Geese, 160
first_calc_done	∼Geese, 164
Model< Array_Type, Data_Counter_Type,	calc_reduced_sequence, 164
Data_Rule_Type, Data_Rule_Dyn_Type >,	calc_sequence, 165
196	colnames, 165
Flock, 150	delete_rengine, 172
~Flock, 151	delete_support, 172
add_data, 151	etype_default, 172
colnames, 152	etype_delatit, 172 etype_duplication, 172
dat, 156	etype_duplication, 172 etype_either, 172
Flock, 151	etype_speciation, 173
get_counters, 152	Geese, 163, 164
get_model, 152	get_annotated_nodes, 165
get_stats_support, 152	get annotations, 165
<del>-</del> · ·	9 = ,
get_stats_target, 152	get_counters, 165
get_support_fun, 153	get_model, 166
init, 153	get_probabilities, 166
initialized, 156	get_rengine, 166
likelihood_joint, 153	get_states, 166
model, 156	get_support_fun, 166
nfunctions, 156	inherit_support, 167
nfuns, 153	init, 167
nleafs, 154	init_node, 167
nnodes, 154	initialized, 173
nterms, 154	likelihood, 167
ntrees, 154	likelihood_exhaust, 167
operator(), 154	map_to_state_id, 173
parse_polytomies, 155	nannotations, 168
print, 155	nfunctions, 173
rengine, 156	nfuns, 168
set_seed, 155	nleafs, 168
support_size, 155	nnodes, 168
flush_data	nodes, 173

nterms, 168	BArrayDense < Cell_Type, Data_Type >, 85
observed_counts, 169	get_col_vec
operator=, 169	BArray< Cell_Type, Data_Type >, 65
parse_polytomies, 169	BArrayDense< Cell_Type, Data_Type >, 85, 86
predict, 169	get_column_major
predict_backend, 170	DEFM, 140
predict_exhaust, 170	get_counters
predict_exhaust_backend, 170	Flock, 152
predict_sim, 170	Geese, 165
print, 170	Model< Array_Type, Data_Counter_Type,
print_nodes, 171	Data_Rule_Type, Data_Rule_Dyn_Type >,
print_observed_counts, 171	183
pset_loc, 173	PhyloCounterData, 215
reduced_sequence, 174	StatsCounter< Array_Type, Data_Type >, 236
sequence, 174	Support< Array_Type, Data_Counter_Type,
set_seed, 171	Data_Rule_Type, Data_Rule_Dyn_Type >,
simulate, 171	242
support_size, 171	get_counts
update_annotations, 171	Support< Array_Type, Data_Counter_Type,
geese, 58	Data_Rule_Type, Data_Rule_Dyn_Type >,
geese-bones.hpp	242
INITIALIZED, 321	get_current_stats
keygen_full, 321	Support< Array_Type, Data_Counter_Type,
RULE_FUNCTION, 321	Data_Rule_Type, Data_Rule_Dyn_Type >,
vec_diff, 321	242
vector_caster, 322	get_data
geese-meat-likelihood.hpp	BArrayDense< Cell_Type, Data_Type >, 86
pset_loop, 323	FreqTable $<$ T $>$ , 159
geese-types.hpp	PowerSet < Array_Type, Data_Rule_Type >, 222
PhyloArray, 329	Support< Array_Type, Data_Counter_Type,
PhyloCounter, 329	Data_Rule_Type, Data_Rule_Dyn_Type >,
PhyloCounters, 329	243
PhyloModel, 329	get_data_ptr
PhyloPowerSet, 329	PowerSet < Array_Type, Data_Rule_Type >, 222
PhyloRule, 330	get_description
PhyloRuleData, 330	Counter< Array_Type, Data_Type >, 131
PhyloRuleDyn, 330	Rule < Array_Type, Data_Type >, 228
PhyloRules, 330	get_descriptions
PhyloRulesDyn, 330	Counters < Array_Type, Data_Type >, 136
PhyloStatsCounter, 330	Rules< Array_Type, Data_Type >, 231
PhyloSupport, 331	StatsCounter< Array_Type, Data_Type >, 236
POS, 328	get_entries
gen_hash	BArray< Cell_Type, Data_Type >, 65
Counters < Array_Type, Data_Type >, 136	BArrayDense < Cell_Type, Data_Type >, 86
gen_key	get_hasher
Model< Array_Type, Data_Counter_Type,	Counter< Array_Type, Data_Type >, 131
Data_Rule_Type, Data_Rule_Dyn_Type >,	get_ID
183	DEFM, 140
get_annotated_nodes	get_index
Geese, 165	FreqTable < T >, 159
get_annotations	get_last_name
Geese, 165	Counting, 24
get_arrays2support	get_likelihoods
Model< 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 >,
183	184
get cell	get_m_order
BArray< Cell Type, Data Type >, 64	DEFM, 140

get_	model DEFM, 141	BArrayDense< Cell_Type, Data_Type >, 86, 87 get_rules
	Flock, 152 Geese, 166	Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,
	n_covars DEFM, 141	186 Support< Array_Type, Data_Counter_Type,
	n_obs DEFM, 141	Data_Rule_Type, Data_Rule_Dyn_Type >, 243
	n_rows DEFM, 141	get_rules_dyn  Model< Array_Type, Data_Counter_Type,  Data_Data_Data_Data_Data_Data_Data_Dat
get_	DEFM, 141	Data_Rule_Type, Data_Rule_Dyn_Type >, 186
get_	name Counter< Array_Type, Data_Type >, 131 Rule< Array_Type, Data_Type >, 228	Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 243
get_	names Counters < Array Type, Data Type >, 137	get_seq Rules< Array Type, Data Type >, 232
	Rules < Array_Type, Data_Type >, 232	get_states
get	StatsCounter< Array_Type, Data_Type >, 236 normalizing_constants	Geese, 166 get_stats_support
<b>-</b>	Model < Array_Type, Data_Counter_Type,	Flock, 152
	Data_Rule_Type, Data_Rule_Dyn_Type >, 184	Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,
get_	parent	186
aet	Node, 208 probabilities	get_stats_support_sizes  Model< Array_Type, Data_Counter_Type,
90'_	Geese, 166	Data_Rule_Type, Data_Rule_Dyn_Type >,
get_	pset  Model< Array_Type, Data_Counter_Type,	186
	Data_Rule_Type, Data_Rule_Dyn_Type >, 184	Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,
get_	pset_arrays  Model< Array_Type, Data_Counter_Type,	187 get stats target
	Data_Rule_Type, Data_Rule_Dyn_Type >, 184	Flock, 152  Model< Array_Type, Data_Counter_Type,
get_	pset_locations  Model< Array_Type, Data_Counter_Type,	Data_Rule_Type, Data_Rule_Dyn_Type >, 187
	Data_Rule_Type, Data_Rule_Dyn_Type >, 184	
get_	pset_probs  Model< Array_Type, Data_Counter_Type,	Geese, 166  Model < Array_Type, Data_Counter_Type,
	Data_Rule_Type, Data_Rule_Dyn_Type >, 185	Data_Rule_Type, Data_Rule_Dyn_Type >, 187
get_	pset_sizes  Model< Array_Type, Data_Counter_Type,	get_X DEFM, 141
	Data_Rule_Type, Data_Rule_Dyn_Type >, 185	get_X_names DEFM, 142
get_	pset_stats	get_Y
	Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 185	DEFM, 142 get_Y_names DEFM, 142
get_	rengine Geese, 166	has_leaf
	Model< Array_Type, Data_Counter_Type,	NodeData, 213
	Data_Rule_Type, Data_Rule_Dyn_Type >,	hasher counters-meat.hpp, 285, 286
get_	186 row_vec	hasher_fun
	BArray< Cell_Type, Data_Type >, 65	Counter< Array_Type, Data_Type >, 133 counters-meat.hpp, 286

hasher_fun_	include/barry/models/defm/defm-meat.hpp, 315
counters-meat.hpp, 289	include/barry/models/defm/defm-types.hpp, 316
Hasher_fun_type	include/barry/models/defm/formula.hpp, 317
typedefs.hpp, 344	include/barry/models/geese.hpp, 319
hashes	include/barry/models/geese/counters.hpp, 312
Support< Array_Type, Data_Counter_Type,	include/barry/models/geese/flock-bones.hpp, 319
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/models/geese/flock-meat.hpp, 320
247	
	include/barry/models/geese/geese-bones.hpp, 320
hashes_initialized	include/barry/models/geese/geese-meat-constructors.hpp,
Support< Array_Type, Data_Counter_Type,	322
Data_Rule_Type, Data_Rule_Dyn_Type >, 247	include/barry/models/geese/geese-meat-likelihood.hpp, 323
:	$include/barry/models/geese/geese-meat-likelihood\_exhaust.hpp,$
i counters most han 200	324
counters-meat.hpp, 289	include/barry/models/geese/geese-meat-predict.hpp,
id	325
Node, 210 idx	include/barry/models/geese/geese-meat-predict_exhaust.hpp,
Phylo rules, 40, 41	325
if	include/barry/models/geese/geese-meat-predict_sim.hpp,
counters-meat.hpp, 286	326
• • •	include/barry/models/geese/geese-meat-simulate.hpp,
IF_MATCHES	326
Counting, 15	include/barry/models/geese/geese-meat.hpp, 327
IF_NOTMATCHES	include/barry/models/geese/geese-node-bones.hpp,
Counting, 15	327
include/barry/barray-bones.hpp, 251	include/barry/models/geese/geese-types.hpp, 328
include/barry/barray-iterator.hpp, 251	include/barry/powerset-bones.hpp, 331
include/barry/barray-meat-operators.hpp, 252	include/barry/powerset-meat.hpp, 332
include/barry/barray-meat.hpp, 256	include/barry/progress.hpp, 332
include/barry/barraycell-bones.hpp, 257	include/barry/rules-bones.hpp, 333
include/barry/barraycell-meat.hpp, 258	include/barry/rules-meat.hpp, 334
include/barry/barraydense-bones.hpp, 258	include/barry/statscounter-bones.hpp, 334
include/barry/barraydense-meat-operators.hpp, 259	include/barry/statscounter-meat.hpp, 335
include/barry/barraydense-meat.hpp, 262	•
include/barry/barraydensecell-bones.hpp, 263	include/barry/support-bones.hpp, 340
include/barry/barraydensecell-meat.hpp, 264	include/barry/support-meat.hpp, 341
include/barry/barraydensecol-bones.hpp, 265	include/barry/typedefs.hpp, 342
include/barry/barraydenserow-bones.hpp, 266	indices
117	NetCounterData, 203
include/barry/barrayrow-bones.hpp, 267	Phylo rules, 45
include/barry/barrayrow-meat.hpp, 267	inherit_support
include/barry/barrayvector-bones.hpp, 269	Geese, 167
include/barry/barrayvector-meat.hpp, 270	init
include/barry/barry-configuration.hpp, 270	Counter< Array_Type, Data_Type >, 131
include/barry/barry-debug.hpp, 272	DEFM, 142
include/barry/barry-macros.hpp, 273	Flock, 153
include/barry/barry.hpp, 274	Geese, 167
include/barry/cell-bones.hpp, 277	Phylo rules, 45
include/barry/cell-meat.hpp, 278	init_fun
include/barry/col-bones.hpp, 278	Counter< Array_Type, Data_Type >, 133
include/barry/counters-bones.hpp, 278	counters-meat.hpp, 287
include/barry/counters-meat.hpp, 279	
include/barry/counters/network-css.hpp, 291	init_fun_ counters-meat hop 290
include/barry/counters/network.hpp, 299	counters-meat.hpp, 290
include/barry/freqtable.hpp, 306	init_node
include/barry/model-bones.hpp, 307	Geese, 167
include/barry/model-meat.hpp, 307	init_support
	PowerSet < Array_Type, Data_Rule_Type >, 222
include/barry/models/defm.hpp, 308	Support< Array_Type, Data_Counter_Type,
include/barry/models/defm/counters.hpp, 309	Data_Rule_Type, Data_Rule_Dyn_Type >,
include/barry/models/defm/defm-bones.hpp, 314	

243	Model < Array_Type, Data_Counter_Type,
INITIALIZED	Data_Rule_Type, Data_Rule_Dyn_Type >,
geese-bones.hpp, 321	187, 188
initialized	likelihood_
Flock, 156	model-meat.hpp, 308
Geese, 173	likelihood_exhaust
insert_cell	Geese, 167
BArray< Cell_Type, Data_Type >, 66	likelihood_joint
BArrayDense < Cell_Type, Data_Type >, 87	Flock, 153
is_col	likelihood_total
BArrayVector< Cell_Type, Data_Type >, 115 BArrayVector_const< Cell_Type, Data_Type >, 119	Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 189
is_dense	logical
BArray< Cell_Type, Data_Type >, 66	Phylo rules, 45
BArrayDense < Cell_Type, Data_Type >, 87	logodds
IS_DUPLICATION	DEFM, 142
Counting, 15	
IS_EITHER	M
Counting, 15	PowerSet < Array_Type, Data_Rule_Type >, 224
is empty	Support< Array_Type, Data_Counter_Type,
BArray< Cell_Type, Data_Type >, 66	Data_Rule_Type, Data_Rule_Dyn_Type >,
BArrayDense< Cell_Type, Data_Type >, 88	247
is leaf	MAKE_DEFM_HASHER
_ Node, 209	Counting, 16
is_motif	MAKE_DUPL_VARS
	Counting, 16
Phylo rules, 45	make_hash
is_row	FreqTable $<$ T $>$ , 159
BArrayVector< Cell_Type, Data_Type >, 115	Мар
BArrayVector_const< Cell_Type, Data_Type >,	barry-configuration.hpp, 272
119	map_to_state_id
IS_SPECIATION	Geese, 173
Counting, 16	MapVec_type
is true	typedefs.hpp, 345
Phylo rules, 41	max_num_elements
iter	Support< Array_Type, Data_Counter_Type,
ConstBArrayRowlter< Cell_Type, Data_Type >,	Data_Rule_Type, Data_Rule_Dyn_Type >,
128	248
120	Model
j	Model < Array_Type, Data_Counter_Type,
counters-meat.hpp, 290	Data_Rule_Type, Data_Rule_Dyn_Type >,
statscounter-meat.hpp, 340	179, 180
• • • • • • • • • • • • • • • • • • • •	model
keygen_defm	Flock, 156
defm-meat.hpp, 316	Model < Array_Type, Data_Counter_Type, Data_Rule_Type
keygen_full	Data_Rule_Dyn_Type >, 175
geese-bones.hpp, 321	$\sim$ Model, 180
keys2support	add_array, 180
Model< Array_Type, Data_Counter_Type,	add_counter, 181
Data_Rule_Type, Data_Rule_Dyn_Type >,	add_hasher, 181
196	add_rule, 181, 182
	add_rule_dyn, 182
lb	arrays2support, 195
PhyloRuleDynData, 218	colnames, 182
likelihood	conditional_prob, 182
Geese, 167	counter_fun, 195
	counters, 195
	delete_counters, 195

delete_rengine, 195	store_psets, 193
delete_rules, 196	support_fun, 200
delete_rules_dyn, 196	support_size, 193
first_calc_done, 196	transform_model, 193
gen_key, 183	transform_model_fun, 201
get_arrays2support, 183	transform_model_term_names, 201
get_counters, 183	update_likelihoods, 194
get_likelihoods, 184	update_normalizing_constants, 194
get_normalizing_constants, 184	update_pset_probs, 194
get_pset, 184	with_pset, 201
get_pset_arrays, 184	model-meat.hpp
get_pset_locations, 184	likelihood_, 308
get_pset_probs, 185	update_normalizing_constant, 308
get_pset_sizes, 185	motif_census
get_pset_stats, 185	DEFM, 143
get_rengine, 186	
get_rules, 186	N
get_rules_dyn, 186	PowerSet < Array_Type, Data_Rule_Type >, 224
get_stats_support, 186	Support< Array_Type, Data_Counter_Type,
get_stats_support_sizes, 186	Data_Rule_Type, Data_Rule_Dyn_Type >,
get_stats_support_sizes_acc, 187	248
get_stats_target, 187	n_counters
get_support_fun, 187	Support< Array_Type, Data_Counter_Type,
keys2support, 196	Data_Rule_Type, Data_Rule_Dyn_Type >,
likelihood, 187, 188	248
likelihood_total, 189	n_free
Model, 179, 180	PowerSet < Array_Type, Data_Rule_Type >, 224
normalizing_constants, 196	n_locked
nrules, 189	PowerSet < Array_Type, Data_Rule_Type >, 224
nrules_dyn, 189	name Counter < Arroy Type Data Type > 100
nterms, 189	Counter< Array_Type, Data_Type >, 133
operator=, 190	counters-meat.hpp, 287
params_last, 197	name_
print, 190	counters-meat.hpp, 290 nannotations
print_stats, 190	Geese, 168
pset_arrays, 197	narray
pset_locations, 197	Node, 210
pset_probs, 197	ncol
pset_sizes, 198	BArray< Cell Type, Data Type >, 67
pset_stats, 198	BArrayDense < Cell Type, Data Type >, 88
rengine, 198	Phylo rules, 41
rules, 198	NET C DATA IDX
rules_dyn, 199	network.hpp, 303
sample, 190, 191	NET_C_DATA_NUM
set_counters, 191	network.hpp, 303
set_rengine, 191 set_rules, 191	NetCounter
set rules dyn, 192	network.hpp, 304
set seed, 192	NetCounterData, 202
set_transform_model, 192	$\sim$ NetCounterData, 203
size, 193	indices, 203
size_unique, 193	NetCounterData, 202
stats_likelihood, 199	numbers, 203
stats_support, 199	NetCounters
stats_support_n_arrays, 199	network.hpp, 304
stats_support_sizes, 200	NetModel
stats_support_sizes_acc, 200	network.hpp, 305
stats_target, 200	NetRule
	network.hpp, 305

NetRules	CSS MATCH TYPE, 294
network.hpp, 305	CSS_NET_COUNTER_LAMBDA_INIT, 294
NetStatsCounter	CSS_PERCEIVED_CELLS, 294
network.hpp, 305	CSS_SIZE, 294
NetSupport	CSS_TRUE_CELLS, 295
network.hpp, 305	network.hpp
Network	BARRY_ZERO_NETWORK, 302
network.hpp, 305	BARRY_ZERO_NETWORK_DENSE, 302
Network counters, 24	NET C DATA IDX, 303
counter_absdiff, 26	NET C DATA NUM, 303
counter ctriads, 27	NetCounter, 304
counter_degree, 27	NetCounters, 304
counter_density, 27	NetModel, 305
counter_diff, 28	NetRule, 305
counter_edges, 28	NetRules, 305
counter_fixed_effect, 28	NetStatsCounter, 305
counter_idegree, 29	NetSupport, 305
counter_idegree15, 29	Network, 305
counter_isolates, 29, 30	NETWORK_COUNTER, 303
counter_istar2, 30	NETWORK COUNTER LAMBDA, 303
counter_logit_intercept, 30	NETWORK RULE, 303
counter_mutual, 30	NETWORK RULE LAMBDA, 304
counter_nodecov, 31	NetworkDense, 306
counter_nodeicov, 31	NETWORKDENSE_COUNTER_LAMBDA, 304
counter_nodematch, 31	rules zerodiag, 306
counter_nodeocov, 31	NETWORK COUNTER
counter_odegree, 31, 32	Network counters, 34
counter_odegree15, 32	network.hpp, 303
counter_ones, 32	NETWORK_COUNTER_LAMBDA
counter ostar2, 33	network.hpp, 303
counter_transition, 33	NETWORK_RULE
counter_transition_formula, 34	network.hpp, 303
counter ttriads, 34	NETWORK RULE LAMBDA
NETWORK COUNTER, 34	network.hpp, 304
rules_dont_become_zero, 34	NetworkData, 203
rules_markov_fixed, 35	~NetworkData, 205
network-css.hpp	directed, 205
counter_css_census01, 295	NetworkData, 204, 205
counter css census02, 295	vertex attr, 205
counter css census03, 295	NetworkDense
counter css census04, 296	network.hpp, 306
counter_css_census05, 296	NETWORKDENSE_COUNTER_LAMBDA
counter_css_census06, 296	network.hpp, 304
counter_css_census07, 296	next
counter css census08, 297	Progress, 226
counter_css_census09, 297	nfunctions
counter css census10, 297	Flock, 156
counter css completely false recip comiss, 297	Geese, 173
counter css completely false recip omiss, 298	nfuns
counter_css_mixed_recip, 298	Flock, 153
counter_css_partially_false_recip_commi, 298	Geese, 168
counter css partially false recip omiss, 299	nleafs
CSS APPEND, 292	Flock, 154
CSS CASE ELSE, 293	Geese, 168
CSS CASE PERCEIVED, 293	nnodes
CSS CASE TRUTH, 293	Flock, 154
CSS CHECK SIZE, 293	Geese, 168
CSS_CHECK_SIZE_INIT, 293	nnozero

D4	
BArray < Cell_Type, Data_Type >, 67	Flock, 154
BArrayDense< Cell_Type, Data_Type >, 88	num
Node, 206	Phylo rules, 41, 42
$\sim$ Node, 208	numbers
annotations, 209	NetCounterData, 203
array, 209	Phylo rules, 46
arrays, 209	•
duplication, 210	obs_start
get_parent, 208	Phylo rules, 46
	observed_counts
id, 210	Geese, 169
is_leaf, 209	offspring
narray, 210	
Node, 207, 208	Node, 210
noffspring, 209	ONE
offspring, 210	CHECK, 56
ord, 210	EXISTS, 58
parent, 211	operator BArrayRow< Cell_Type, Data_Type >
probability, 211	BArrayRow< Cell_Type, Data_Type >, 110
subtree_prob, 211	operator BArrayRow_const< Cell_Type, Data_Type >
visited, 211	BArrayRow_const < Cell_Type, Data_Type >, 112
NodeData, 212	operator Cell Type
	BArrayCell< Cell_Type, Data_Type >, 75
blengths, 212	BArrayCell_const< Cell_Type, Data_Type >, 77
duplication, 213	BArrayDenseCell< Cell_Type, Data_Type >, 97
has_leaf, 213	Cell< Cell_Type >, 124
NodeData, 212	
states, 213	operator std::vector < Cell_Type >
nodes	BArrayVector< Cell_Type, Data_Type >, 116
Geese, 173	BArrayVector_const< Cell_Type, Data_Type >,
noexcept	119
counters-meat.hpp, 290	operator!=
noffspring	BArrayCell_const< Cell_Type, Data_Type >, 77
Node, 209	BArrayRow_const< Cell_Type, Data_Type >, 112
NONE	BArrayVector_const< Cell_Type, Data_Type >,
CHECK, 56	119
EXISTS, 58	Cell< Cell_Type >, 124
normalizing_constants	operator<
•—	BArrayCell_const< Cell_Type, Data_Type >, 77
	BArrayRow_const< Cell_Type, Data_Type >, 112
Data_Rule_Type, Data_Rule_Dyn_Type >,	BArrayVector_const< Cell_Type, Data_Type >,
196	120
nrow	operator<=
BArray < Cell_Type, Data_Type >, 67	BArrayCell_const< Cell_Type, Data_Type >, 78
BArrayDense< Cell_Type, Data_Type >, 88	
Phylo rules, 41	BArrayRow_const< Cell_Type, Data_Type >, 113
nrules	BArrayVector_const< Cell_Type, Data_Type >,
Model < Array_Type, Data_Counter_Type,	120
Data_Rule_Type, Data_Rule_Dyn_Type >,	operator>
189	BArrayCell_const< Cell_Type, Data_Type >, 78
nrules_dyn	BArrayRow_const< Cell_Type, Data_Type >, 113
Model< Array_Type, Data_Counter_Type,	BArrayVector_const< Cell_Type, Data_Type >,
Data_Rule_Type, Data_Rule_Dyn_Type >,	120
189	operator>=
nterms	BArrayCell_const< Cell_Type, Data_Type >, 78
	BArrayRow_const< Cell_Type, Data_Type >, 113
Flock, 154	BArrayVector_const< Cell_Type, Data_Type >,
Geese, 168	120
Model Array_Type, Data_Counter_Type,	operator*=
Data_Rule_Type, Data_Rule_Dyn_Type >,	BArray< Cell_Type, Data_Type >, 67
189	
ntrees	BArrayDense Coll Tipe, Data Tipe >, 75
	BArrayDense < Cell_Type, Data_Type >, 89

BArrayDenseCell< Cell_Type, Data_Type >, 97 BArrayRow< Cell_Type, Data_Type >, 110 BArrayVector< Cell_Type, Data_Type >, 116 operator() BArray< Cell_Type, Data_Type >, 67 barray-meat-operators.hpp, 255 BArrayDense< Cell_Type, Data_Type >, 88, 89 BArrayDenseCol< Cell_Type, Data_Type >, 101	BArrayCell< Cell_Type, Data_Type >, 76 BArrayCell_const< Cell_Type, Data_Type >, 78 BArrayDense< Cell_Type, Data_Type >, 91 BArrayDenseCell< Cell_Type, Data_Type >, 99 BArrayRow< Cell_Type, Data_Type >, 111 BArrayRow_const< Cell_Type, Data_Type >, 113 BArrayVector< Cell_Type, Data_Type >, 117 BArrayVector_const< Cell_Type, Data_Type >, 2000
BArrayDenseCol_const< Cell_Type, Data_Type >, 103	120 Cell< Cell_Type >, 125
BArrayDenseRow< Cell_Type, Data_Type >, 106	operator[]
BArrayDenseRow_const< Cell_Type, Data_Type >, 108	Counters< Array_Type, Data_Type >, 138 PhyloCounterData, 215
Flock, 154	PowerSet < Array_Type, Data_Rule_Type >, 222
Phylo rules, 42	ord
PhyloCounterData, 215	Node, 210
PhyloRuleDynData, 217	out_of_range
Rule< Array_Type, Data_Type >, 229 Rules< Array_Type, Data_Type >, 232	BArray< Cell_Type, Data_Type >, 69 BArrayDense< Cell_Type, Data_Type >, 91
vecHasher< T >, 249	Britay Bense < Gen_Type, Bata_Type >, T
operator+=	params_last
BArray< Cell_Type, Data_Type >, 68	Model< Array_Type, Data_Counter_Type,
BArrayCell< Cell_Type, Data_Type >, 75	Data_Rule_Type, Data_Rule_Dyn_Type >,
BArrayDense< Cell_Type, Data_Type >, 89	197
BArrayDenseCell< Cell_Type, Data_Type >, 98	parent
BArrayRow< Cell_Type, Data_Type >, 110	Node, 211
BArrayVector< Cell_Type, Data_Type >, 116	parse_polytomies
operator-=	Flock, 155
BArray< Cell_Type, Data_Type >, 68	Geese, 169
BArrayCell< Cell_Type, Data_Type >, 75	Phylo counters, 47
BArrayDense < Cell_Type, Data_Type >, 90	counter_co_opt, 48
BArrayDenseCell< Cell_Type, Data_Type >, 98	counter_cogain, 49
BArrayRow< Cell_Type, Data_Type >, 110	counter_gains, 49
BArrayVector< Cell_Type, Data_Type >, 116	counter_gains_from_0, 49
operator/=	counter_gains_k_offspring, 49
BArray< Cell_Type, Data_Type >, 69	counter_genes_changing, 50
BArrayCell< Cell_Type, Data_Type >, 75	counter_k_genes_changing, 50
BArrayDense < Cell_Type, Data_Type >, 90	counter_less_than_p_prop_genes_changing, 50
BArrayDenseCell< Cell_Type, Data_Type >, 98	counter_longest, 50
BArrayRow< Cell_Type, Data_Type >, 111	counter_loss, 51
BArrayVector< Cell_Type, Data_Type >, 116	counter_maxfuns, 51
operator=	counter_neofun, 51
BArray< Cell_Type, Data_Type >, 69	counter_neofun_a2b, 51 counter_overall_changes, 52
BArrayCell< Cell_Type, Data_Type >, 75	counter_overall_gains, 52
BArrayDense < Cell_Type, Data_Type >, 90, 91	counter_overall_gains_from_0, 52
BArrayDenseCell < Cell_Type, Data_Type >, 98	counter_overall_loss, 52
BArrayRow< Cell_Type, Data_Type >, 111	counter pairwise first gain, 53
BArrayVector < Cell_Type, Data_Type >, 117	counter_pairwise_neofun_singlefun, 53
Cell< Cell_Type >, 124, 125	counter_pairwise_overall_change, 53
Counter< Array_Type, Data_Type >, 131, 132 Counters< Array_Type, Data_Type >, 137	counter_pairwise_preserving, 53
Geese, 169	counter_preserve_pseudogene, 54
Model< Array_Type, Data_Counter_Type,	counter_prop_genes_changing, 54
Data_Rule_Type, Data_Rule_Dyn_Type >,	counter_subfun, 54
190	Phylo rules, 35
Rules < Array_Type, Data_Type >, 233	~DEFMCounterData, 43
operator==	∼DEFMData, 43
BArray< Cell_Type, Data_Type >, 69	$\sim$ DEFMRuleDynData, 43
	array, 44

at, 39	shrink_to_fit, 216
counts, 44	size, 216
covar_sort, 44	PhyloCounters
covar_used, 44	geese-types.hpp, 329
covariates, 44	PhyloModel
DEFMCounter, 37	geese-types.hpp, 329
DEFMCounterData, 39	PhyloPowerSet
DEFMCounters, 37	geese-types.hpp, 329
DEFMData, 39	PhyloRule
DEFMModel, 37	geese-types.hpp, 330
DEFMRule, 38	PhyloRuleData
DEFMRuleData, 40	geese-types.hpp, 330
DEFMRuleDyn, 38	PhyloRuleDyn
DEFMRuleDynData, 40	geese-types.hpp, 330
DEFMRules, 38	PhyloRuleDynData, 216
DEFMRulesDyn, 38	~PhyloRuleDynData, 217
DEFMStatsCounter, 38	counts, 218
DEFMSupport, 38	duplication, 218
idx, 40, 41	lb, 218
indices, 45	operator(), 217
init, 45	PhyloRuleDynData, 217
is_motif, 45	pos, 218
is_true, 41	ub, 218
logical, 45	PhyloRules
ncol, 41	geese-types.hpp, 330
nrow, 41	PhyloRulesDyn
num, 41, 42	geese-types.hpp, 330
numbers, 46	PhyloStatsCounter
obs_start, 46	geese-types.hpp, 330
operator(), 42	PhyloSupport 201
print, 42	geese-types.hpp, 331
rule_dyn_limit_changes, 42	POS
rule_leafs, 43	barraydense-meat-operators.hpp, 260
X_ncol, 46	barraydense-meat.hpp, 262
X_nrow, 46	barraydensecell-bones.hpp, 264
PHYLO_CHECK_MISSING	barraydensecell-meat.hpp, 264
Counting, 16	barraydensecol-bones.hpp, 265
PHYLO_COUNTER_LAMBDA	barraydenserow-bones.hpp, 267
Counting, 17	geese-types.hpp, 328
PHYLO_RULE_DYN_LAMBDA	pos Phylo Pylo Dyro Poto 210
Counting, 17 PHYLO_RULE_LAMBDA	PhyloRuleDynData, 218 POS N
Counting, 17	barraydense-meat-operators.hpp, 260
PhyloArray	barraydense-meat.hpp, 263
	barraydensecol-bones.hpp, 265
geese-types.hpp, 329 PhyloCounter	barraydenserow-bones.hpp, 267
geese-types.hpp, 329	PowerSet
PhyloCounterData, 213	PowerSet < Array_Type, Data_Rule_Type >, 220
at, 214	PowerSet < Array_Type, Data_Rule_Type >, 219
begin, 215	~PowerSet, 221
empty, 215	add_rule, 221
end, 215	begin, 221
get_counters, 215	calc, 221
operator(), 215	coordinates_free, 223
operator(), 215	coordinates_locked, 223
PhyloCounterData, 214	data, 223
push_back, 216	EmptyArray, 224
reserve, 216	end, 222
.000, 70, 270	Orio, ELL

get_data, 222	pset_arrays
get_data_ptr, 222	Model < Array_Type, Data_Counter_Type,
init_support, 222	Data_Rule_Type, Data_Rule_Dyn_Type >,
M, 224	197
N, 224	pset_loc
	• —
n_free, 224	Geese, 173
n_locked, 224	pset_locations
operator[], 222	Model < Array_Type, Data_Counter_Type,
PowerSet, 220	Data_Rule_Type, Data_Rule_Dyn_Type >,
reset, 223	197
rules, 224	pset_loop
rules_deleted, 225	geese-meat-likelihood.hpp, 323
size, 223	pset_probs
predict	Model < Array_Type, Data_Counter_Type,
Geese, 169	Data_Rule_Type, Data_Rule_Dyn_Type >,
predict_backend	197
Geese, 170	pset_sizes
predict_exhaust	Model < Array_Type, Data_Counter_Type,
Geese, 170	Data_Rule_Type, Data_Rule_Dyn_Type >,
predict_exhaust_backend	198
Geese, 170	pset_stats
predict_sim	Model < Array_Type, Data_Counter_Type,
Geese, 170	Data_Rule_Type, Data_Rule_Dyn_Type >,
print	198
BArray< Cell_Type, Data_Type >, 69	push_back
BArrayDense< Cell_Type, Data_Type >, 91	PhyloCounterData, 216
DEFM, 143	ThylodountorBata, 210
	README.md, 347
Flock, 155	
FreqTable $<$ T $>$ , 159	reduced_sequence
Geese, 170	Geese, 174
Geese, 170  Model< Array Type. Data Counter Type.	Geese, 174 rengine
Model < Array_Type, Data_Counter_Type,	rengine
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,	rengine Flock, 156
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 190	rengine Flock, 156 Model< Array_Type, Data_Counter_Type,
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 190 Phylo rules, 42	rengine Flock, 156 Model< 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 >, 190	rengine Flock, 156 Model< Array_Type, Data_Counter_Type,
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 190 Phylo rules, 42 Support< Array_Type, Data_Counter_Type,	rengine Flock, 156 Model< 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 >, 190 Phylo rules, 42 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 190  Phylo rules, 42  Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 244	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 190 Phylo rules, 42 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 244 print_n	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290 reserve
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 190 Phylo rules, 42 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 244  print_n BArray< Cell_Type, Data_Type >, 70	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray< Cell_Type, Data_Type >, 70
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 190 Phylo rules, 42 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 244 print_n	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 91
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 190 Phylo rules, 42 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 244  print_n BArray< Cell_Type, Data_Type >, 70	rengine Flock, 156 Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray < Cell_Type, Data_Type >, 70 BArrayDense < Cell_Type, Data_Type >, 91 FreqTable < T >, 159
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 190 Phylo rules, 42 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 244  print_n BArray< Cell_Type, Data_Type >, 70  print_nodes Geese, 171	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 91
Model< Array_Type, Data_Counter_Type,	rengine Flock, 156 Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray < Cell_Type, Data_Type >, 70 BArrayDense < Cell_Type, Data_Type >, 91 FreqTable < T >, 159
Model< Array_Type, Data_Counter_Type,	rengine Flock, 156 Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray < Cell_Type, Data_Type >, 70 BArrayDense < Cell_Type, Data_Type >, 91 FreqTable < T >, 159 PhyloCounterData, 216  reset
Model< Array_Type, Data_Counter_Type,	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 91 FreqTable< T >, 159 PhyloCounterData, 216  reset PowerSet< Array_Type, Data_Rule_Type >, 223
Model< Array_Type, Data_Counter_Type,         Data_Rule_Type, Data_Rule_Dyn_Type >,         190  Phylo rules, 42  Support< Array_Type, Data_Counter_Type,         Data_Rule_Type, Data_Rule_Dyn_Type >,         244  print_n         BArray< Cell_Type, Data_Type >, 70  print_nodes         Geese, 171  print_observed_counts         Geese, 171  print_stats         Model< Array_Type, Data_Counter_Type,	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 91 FreqTable< T >, 159 PhyloCounterData, 216  reset PowerSet< Array_Type, Data_Rule_Type >, 223  reset_array
Model< Array_Type, Data_Counter_Type,	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 91 FreqTable< T >, 159 PhyloCounterData, 216  reset PowerSet< Array_Type, Data_Rule_Type >, 223  reset_array StatsCounter< Array_Type, Data_Type >, 236
Model< Array_Type, Data_Counter_Type,         Data_Rule_Type, Data_Rule_Dyn_Type >,         190  Phylo rules, 42  Support< Array_Type, Data_Counter_Type,         Data_Rule_Type, Data_Rule_Dyn_Type >,         244  print_n         BArray< Cell_Type, Data_Type >, 70  print_nodes         Geese, 171  print_observed_counts         Geese, 171  print_stats         Model< Array_Type, Data_Counter_Type,	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 91 FreqTable< T >, 159 PhyloCounterData, 216  reset PowerSet< Array_Type, Data_Rule_Type >, 223  reset_array StatsCounter< Array_Type, Data_Type >, 236 Support< Array_Type, Data_Counter_Type,
Model< Array_Type, Data_Counter_Type,         Data_Rule_Type, Data_Rule_Dyn_Type >,         190 Phylo rules, 42 Support< Array_Type, Data_Counter_Type,         Data_Rule_Type, Data_Rule_Dyn_Type >,         244  print_n         BArray< Cell_Type, Data_Type >, 70  print_nodes         Geese, 171  print_observed_counts         Geese, 171  print_stats         Model< Array_Type, Data_Counter_Type,         Data_Rule_Type, Data_Rule_Dyn_Type >,         190	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 91 FreqTable< T >, 159 PhyloCounterData, 216  reset PowerSet< Array_Type, Data_Rule_Type >, 223  reset_array StatsCounter< Array_Type, Data_Type >, 236
Model< Array_Type, Data_Counter_Type,         Data_Rule_Type, Data_Rule_Dyn_Type >,         190 Phylo rules, 42 Support< Array_Type, Data_Counter_Type,         Data_Rule_Type, Data_Rule_Dyn_Type >,         244  print_n         BArray< Cell_Type, Data_Type >, 70  print_nodes         Geese, 171  print_observed_counts         Geese, 171  print_stats         Model< Array_Type, Data_Counter_Type,         Data_Rule_Type, Data_Rule_Dyn_Type >,         190  printf_barry	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 91 FreqTable< T >, 159 PhyloCounterData, 216  reset PowerSet< Array_Type, Data_Rule_Type >, 223  reset_array StatsCounter< Array_Type, Data_Type >, 236 Support< Array_Type, Data_Counter_Type,
Model< Array_Type, Data_Counter_Type,	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 91 FreqTable< T >, 159 PhyloCounterData, 216  reset PowerSet< Array_Type, Data_Rule_Type >, 223  reset_array StatsCounter< Array_Type, Data_Type >, 236 Support< Array_Type, Data_Rule_Dyn_Type, >, 244
Model< Array_Type, Data_Counter_Type,	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 91 FreqTable< T >, 159 PhyloCounterData, 216  reset PowerSet< Array_Type, Data_Rule_Type >, 223  reset_array StatsCounter< Array_Type, Data_Type >, 236 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 244  resize
Model< Array_Type, Data_Counter_Type,	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 91 FreqTable< T >, 159 PhyloCounterData, 216  reset PowerSet< Array_Type, Data_Rule_Type >, 223  reset_array StatsCounter< Array_Type, Data_Type >, 236 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 244  resize BArray< Cell_Type, Data_Type >, 70
Model< Array_Type, Data_Counter_Type,	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 91 FreqTable< T >, 159 PhyloCounterData, 216  reset PowerSet< Array_Type, Data_Rule_Type >, 223  reset_array StatsCounter< Array_Type, Data_Type >, 236 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 244  resize BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 92
Model< Array_Type, Data_Counter_Type,	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 91 FreqTable< T >, 159 PhyloCounterData, 216  reset PowerSet< Array_Type, Data_Rule_Type >, 223  reset_array StatsCounter< Array_Type, Data_Type >, 236 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 244  resize BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 92 Entries< Cell_Type >, 149
Model< Array_Type, Data_Counter_Type,	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 91 FreqTable< T >, 159 PhyloCounterData, 216  reset PowerSet< Array_Type, Data_Rule_Type >, 223  reset_array StatsCounter< Array_Type, Data_Type >, 236 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 244  resize BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 92
Model< Array_Type, Data_Counter_Type,	rengine Flock, 156 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 91 FreqTable< T >, 159 PhyloCounterData, 216  reset PowerSet< Array_Type, Data_Rule_Type >, 223  reset_array StatsCounter< Array_Type, Data_Type >, 236 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 244  resize BArray< Cell_Type, Data_Type >, 70 BArrayDense< Cell_Type, Data_Type >, 92 Entries< Cell_Type >, 149
Model< Array_Type, Data_Counter_Type,	rengine Flock, 156 Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray < Cell_Type, Data_Type >, 70 BArrayDense < Cell_Type, Data_Type >, 91 FreqTable < T >, 159 PhyloCounterData, 216  reset PowerSet < Array_Type, Data_Rule_Type >, 223  reset_array StatsCounter < Array_Type, Data_Type >, 236 Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 244  resize BArray < Cell_Type, Data_Type >, 70 BArrayDense < Cell_Type, Data_Type >, 92 Entries < Cell_Type >, 149 statscounter-meat.hpp, 337  return
Model< Array_Type, Data_Counter_Type,	rengine Flock, 156 Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray < Cell_Type, Data_Type >, 70 BArrayDense < Cell_Type, Data_Type >, 91 FreqTable < T >, 159 PhyloCounterData, 216  reset PowerSet < Array_Type, Data_Rule_Type >, 223  reset_array StatsCounter < Array_Type, Data_Type >, 236 Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 244  resize BArray < Cell_Type, Data_Type >, 70 BArrayDense < Cell_Type, Data_Type >, 92 Entries < Cell_Type >, 149 statscounter-meat.hpp, 337  return counters-meat.hpp, 291
Model< Array_Type, Data_Counter_Type,	rengine Flock, 156 Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray < Cell_Type, Data_Type >, 70 BArrayDense < Cell_Type, Data_Type >, 91 FreqTable < T >, 159 PhyloCounterData, 216  reset PowerSet < Array_Type, Data_Rule_Type >, 223  reset_array StatsCounter < Array_Type, Data_Type >, 236 Support < Array_Type, Data_Rule_Dyn_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 244  resize BArray < Cell_Type, Data_Type >, 70 BArrayDense < Cell_Type, Data_Type >, 92 Entries < Cell_Type >, 149 statscounter-meat.hpp, 337  return counters-meat.hpp, 291 statscounter-meat.hpp, 340
Model< Array_Type, Data_Counter_Type,	rengine Flock, 156 Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198  res counters-meat.hpp, 290  reserve BArray < Cell_Type, Data_Type >, 70 BArrayDense < Cell_Type, Data_Type >, 91 FreqTable < T >, 159 PhyloCounterData, 216  reset PowerSet < Array_Type, Data_Rule_Type >, 223  reset_array StatsCounter < Array_Type, Data_Type >, 236 Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 244  resize BArray < Cell_Type, Data_Type >, 70 BArrayDense < Cell_Type, Data_Type >, 92 Entries < Cell_Type >, 149 statscounter-meat.hpp, 337  return counters-meat.hpp, 291

rm_cell	rules_deleted
BArray< Cell_Type, Data_Type >, 70	PowerSet < Array_Type, Data_Rule_Type >, 225
BArrayDense < Cell_Type, Data_Type >, 92	rules_dont_become_zero
ROW	Network counters, 34
barray-meat-operators.hpp, 253	rules_dyn
barray-meat.hpp, 257	Model < Array_Type, Data_Counter_Type,
barraydense-meat-operators.hpp, 260	Data_Rule_Type, Data_Rule_Dyn_Type >,
barraydense-meat.hpp, 263	199
row	rules_markov_fixed
BArray< Cell_Type, Data_Type >, 70	Network counters, 35
BArrayDense< Cell_Type, Data_Type >, 92	rules_zerodiag
Row type	network.hpp, 306
typedefs.hpp, 345	• • •
rowsum	sample
BArrayDense < Cell_Type, Data_Type >, 92	Model< Array_Type, Data_Counter_Type,
Rule	Data_Rule_Type, Data_Rule_Dyn_Type >,
Rule < Array_Type, Data_Type >, 227	190, 191
Rule < Array_Type, Data_Type >, 226	sequence
∼Rule, 227	Geese, 174
D, 228	set_counters
get_description, 228	Model< Array_Type, Data_Counter_Type,
get_name, 228	Data_Rule_Type, Data_Rule_Dyn_Type >,
operator(), 229	191
Rule, 227	StatsCounter< Array_Type, Data_Type >, 237
rule_dyn_limit_changes	Support< Array_Type, Data_Counter_Type,
Phylo rules, 42	Data_Rule_Type, Data_Rule_Dyn_Type >,
rule_fun_default	244
rules-bones.hpp, 333	set_data
Rule_fun_type	BArray< Cell_Type, Data_Type >, 71
typedefs.hpp, 345	BArrayDense < Cell_Type, Data_Type >, 93
RULE_FUNCTION	set_hasher
barry.hpp, 277	Counter< Array_Type, Data_Type >, 132
geese-bones.hpp, 321	set_names
RULE LAMBDA	DEFM, 143
barry.hpp, 277	set_rengine
rule_leafs	Model < Array_Type, Data_Counter_Type,
Phylo rules, 43	Data_Rule_Type, Data_Rule_Dyn_Type >,
Rules	191
Rules< Array_Type, Data_Type >, 230	set_rules
rules	Model < 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 >,	191
198	Support< Array_Type, Data_Counter_Type,
PowerSet < Array_Type, Data_Rule_Type >, 224	Data_Rule_Type, Data_Rule_Dyn_Type >,
Rules< Array_Type, Data_Type >, 229	245
~Rules, 230	set_rules_dyn
add rule, 231	Model < Array_Type, Data_Counter_Type,
begin, 231	Data_Rule_Type, Data_Rule_Dyn_Type >,
end, 231	192
get_descriptions, 231	Support< Array_Type, Data_Counter_Type,
get_names, 232	Data_Rule_Type, Data_Rule_Dyn_Type >,
get_seq, 232	245
	set seed
operator, 233	Flock, 155
operator=, 233	Geese, 171
Rules, 230	Model< Array_Type, Data_Counter_Type,
size, 233	Data_Rule_Type, Data_Rule_Dyn_Type >,
rules-bones.hpp	192
rule_fun_default, 333	set_transform_model

Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 192	stats_target  Model< Array_Type, Data_Counter_Type,  Data_Rule_Type, Data_Rule_Dyn_Type >,
shrink_to_fit	200
PhyloCounterData, 216	StatsCounter
simulate	StatsCounter< Array_Type, Data_Type >, 234,
DEFM, 143	235
Geese, 171	StatsCounter< Array_Type, Data_Type >, 233
size	∼StatsCounter, 235
BArrayDenseCol< Cell_Type, Data_Type >, 101	add_counter, 235
BArrayDenseCol_const< Cell_Type, Data_Type >,	count_all, 235
104	count_current, 236
BArrayDenseRow< Cell_Type, Data_Type >, 106	count_init, 236
BArrayDenseRow_const< Cell_Type, Data_Type	get_counters, 236
>, 108	get_descriptions, 236
BArrayVector< Cell_Type, Data_Type >, 117	get_names, 236
BArrayVector_const< Cell_Type, Data_Type >,	reset_array, 236
121	set_counters, 237
Counters< Array_Type, Data_Type >, 138	size, 237
FreqTable < T >, 160  Model < Array Type. Data Counter Type.	StatsCounter, 234, 235
	statscounter-meat.hpp clear, 336
Data_Rule_Type, Data_Rule_Dyn_Type >, 193	counter, 338
PhyloCounterData, 216	counter_deleted, 339
PowerSet< Array_Type, Data_Rule_Type >, 223	counters, 339
Rules < Array_Type, Data_Type >, 233	counters_, 339
StatsCounter< Array_Type, Data_Type >, 237	current_stats, 339
size_unique	EmptyArray, 339
Model< Array_Type, Data_Counter_Type,	f_, 340
Data_Rule_Type, Data_Rule_Dyn_Type >,	for, 336
193	j, 340
sort_array	resize, 337
typedefs.hpp, 345	return, 340
source	STATSCOUNTER_TEMPLATE, 336-338
Entries< Cell_Type >, 149	STATSCOUNTER_TEMPLATE_ARGS, 336
states	STATSCOUNTER_TYPE, 336
NodeData, 213	STATSCOUNTER_TEMPLATE
Statistical Models, 24	statscounter-meat.hpp, 336–338
stats_likelihood	STATSCOUNTER_TEMPLATE_ARGS
Model < Array_Type, Data_Counter_Type,	statscounter-meat.hpp, 336
Data_Rule_Type, Data_Rule_Dyn_Type >, 199	STATSCOUNTER_TYPE statscounter-meat.hpp, 336
stats support	store_psets
Model 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 >,
199	193
stats_support_n_arrays	subtree_prob
Model< Array_Type, Data_Counter_Type,	Node, 211
Data_Rule_Type, Data_Rule_Dyn_Type >,	Support
199	Support< Array_Type, Data_Counter_Type,
stats_support_sizes	Data_Rule_Type, Data_Rule_Dyn_Type >,
Model< Array_Type, Data_Counter_Type,	239, 240
Data_Rule_Type, Data_Rule_Dyn_Type >,	
200	Data_Rule_Dyn_Type >, 237
stats_support_sizes_acc	~Support, 240
Model Array_Type, Data_Counter_Type,	add_counter, 240
Data_Rule_Type, Data_Rule_Dyn_Type >,	add_rule, 240, 241
200	add_rule_dyn, <mark>24</mark> 1

calc, 241 change stats, 245	BArray< Cell_Type, Data_Type >, 72 BArrayDense< Cell_Type, Data_Type >, 94
coordiantes_n_free, 245	toggle_lock
coordiantes_n_locked, 246	BArray< Cell_Type, Data_Type >, 72
coordinates_free, 246	BArrayDense< Cell_Type, Data_Type >, 94
coordinates locked, 246	transform_model
current_stats, 246	Model < Array_Type, Data_Counter_Type,
delete_counters, 246	Data_Rule_Type, Data_Rule_Dyn_Type >,
delete rules, 247	193
delete rules dyn, 247	transform_model_fun
eval rules dyn, 242	Model< Array_Type, Data_Counter_Type,
get_counters, 242	Data_Rule_Type, Data_Rule_Dyn_Type >,
get_counts, 242	201
get_current_stats, 242	transform_model_term_names
get_data, 243	Model< Array_Type, Data_Counter_Type,
get_rules, 243	Data_Rule_Type, Data_Rule_Dyn_Type >,
get_rules_dyn, 243	201
hashes, 247	transpose
hashes_initialized, 247	BArray< Cell_Type, Data_Type >, 72
init_support, 243	BArrayDense < Cell_Type, Data_Type >, 72
M, 247	TWO
max_num_elements, 248	CHECK, 56
N, 248	EXISTS, 58
n_counters, 248	typedefs.hpp
print, 244	Col_type, 344
reset_array, 244	Counter_fun_type, 344
set_counters, 244	
set_rules, 245	Counts_type, 344
	Hasher_fun_type, 344
set_rules_dyn, 245	MapVec_type, 345
Support, 239, 240	Row_type, 345
support-meat.hpp	Rule_fun_type, 345
BARRY_SUPPORT_MEAT_HPP, 341	sort_array, 345
support_fun  Madel C Arrey Type Date Counter Type	vec_equal, 346
Model < Array_Type, Data_Counter_Type,	vec_equal_approx, 346
Data_Rule_Type, Data_Rule_Dyn_Type >,	vec_inner_prod, 346, 347
200	ub
support_size	PhyloRuleDynData, 218
Flock, 155	UKNOWN
Geese, 171	EXISTS, 58
Model < Array_Type, Data_Counter_Type,	UNI_SUB
Data_Rule_Type, Data_Rule_Dyn_Type >,	counters.hpp, 312
193	update_annotations
swap_cells	Geese, 171
BArray Cell_Type, Data_Type >, 71	update_likelihoods
BArrayDense< Cell_Type, Data_Type >, 93	Model Array_Type, Data_Counter_Type,
swap_cols	Data_Rule_Type, Data_Bule_Dyn_Type >,
BArray Cell_Type, Data_Type >, 71	194
BArrayDense< Cell_Type, Data_Type >, 93	update_normalizing_constant
swap_rows	model-meat.hpp, 308
BArray Cell_Type, Data_Type >, 71	update_normalizing_constants
BArrayDense< Cell_Type, Data_Type >, 93	
target	
Entries < Cell_Type >, 149	Data_Rule_Type, Data_Rule_Dyn_Type >,
this	194
barray-meat-operators.hpp, 256	update_pset_probs
TMP_HASHER_CALL	Model < Array_Type, Data_Counter_Type,
counters-meat.hpp, 281	Data_Rule_Type, Data_Rule_Dyn_Type >, 194
toggle_cell	104

```
val
     Entries < Cell_Type >, 149
value
     Cell< Cell_Type >, 125
vec_diff
    geese-bones.hpp, 321
vec equal
     typedefs.hpp, 346
vec_equal_approx
     typedefs.hpp, 346
vec_inner_prod
     typedefs.hpp, 346, 347
vecHasher< T >, 248
    operator(), 249
vector_caster
    geese-bones.hpp, 322
vertex_attr
    NetworkData, 205
visited
     BArray< Cell_Type, Data_Type >, 73
     BArrayDense < Cell_Type, Data_Type >, 96
     Cell< Cell_Type >, 125
    Node, 211
with pset
     Model<
                 Array_Type,
                                  Data_Counter_Type,
         Data_Rule_Type, Data_Rule_Dyn_Type >,
         201
X_ncol
     Phylo rules, 46
X nrow
     Phylo rules, 46
ZERO CELL
    barraydense-meat.hpp, 263
    barraydensecol-bones.hpp, 266
    barraydenserow-bones.hpp, 267
zero col
     {\sf BArray}{<}\,{\sf Cell\_Type},\,{\sf Data\_Type}>, {\sf \color{red}{72}}
     BArrayDense < Cell_Type, Data_Type >, 94
     BArray< Cell_Type, Data_Type >, 72
     BArrayDense < Cell_Type, Data_Type >, 94
```