barry: Your go-to motif accountant 0.0-1

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1 Main Page	1
2 Module Index	5
2.1 Modules	5
3 Hierarchical Index	7
3.1 Class Hierarchy	7
4 Class Index	9
4.1 Class List	9
5 File Index	11
5.1 File List	11
6 Module Documentation	13
6.1 Counting	13
6.1.1 Detailed Description	15
6.1.2 Macro Definition Documentation	15
6.1.2.1 IF_MATCHES	15
6.1.2.2 IF_NOTMATCHES	15
6.1.2.3 IS_DUPLICATION	15
6.1.2.4 IS_EITHER	16
6.1.2.5 IS_SPECIATION	16
6.1.2.6 MAKE_DEFM_HASHER	16
6.1.2.7 MAKE_DUPL_VARS	
6.1.2.8 PHYLO_CHECK_MISSING	17
6.1.2.9 PHYLO_COUNTER_LAMBDA	
6.1.2.10 PHYLO RULE DYN LAMBDA	
6.1.3 Function Documentation	17
6.1.3.1 counter_co_opt()	
6.1.3.2 counter_cogain()	
6.1.3.3 counter_gains()	
6.1.3.4 counter_gains_from_0()	
6.1.3.5 counter_gains_k_offspring()	
6.1.3.6 counter_genes_changing()	
6.1.3.7 counter_k_genes_changing()	
6.1.3.8 counter_less_than_p_prop_genes_changing()	
6.1.3.9 counter_longest()	
6.1.3.10 counter_loss()	
6.1.3.11 counter_maxfuns()	
6.1.3.12 counter_neofun()	
6.1.3.13 counter_neofun_a2b()	
6.1.3.14 counter_overall_changes()	
6.1.3.15 counter_overall_gains()	
o.n.o.ro counter_overan_game()	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
6.3.2.2 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
6.3.2.12 counter_idegree15() [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 counter_istar2()</b> [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 counter_odegree()</b> [1/2]	32
<b>6.3.2.24 counter_odegree()</b> [2/2]	32
<b>6.3.2.25</b> counter_odegree15() [1/2]	32
6.3.2.26 counter_odegree15() [2/2]	32
6.3.2.27 counter_ones()	32

6.3.2.28 counter_ostar2() [1/2]	 . 33
6.3.2.29 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	 . 39
6.4.3.1 at()	 . 39
<b>6.4.3.2 DEFMCounterData()</b> [1/2]	 . 39
<b>6.4.3.3 DEFMCounterData()</b> [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

43
43
43
44
44
44
44
44
44
45
45
45
45
45
45
46
46
46
46
46
47
48
48
48
49
49
49
50
50
50
50
51
51
51
51
52
52
52
52
53
53
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	55
7.1.1 Detailed Description	55
7.2 barry::counters Namespace Reference	55
7.2.1 Detailed Description	55
7.3 barry::counters::network Namespace Reference	56
7.4 CHECK Namespace Reference	56
7.4.1 Detailed Description	56
7.4.2 Variable Documentation	56
7.4.2.1 BOTH	56
7.4.2.2 NONE	56
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
8.1.2.4 BArray() [4/6]	63
<b>8.1.2.5 BArray()</b> [5/6]	63
<b>8.1.2.6 BArray()</b> [6/6]	63
8.1.2.7 ~BArray()	63

8.1.3 N	Member Function Documentation	63
	8.1.3.1 clear()	63
	8.1.3.2 col()	64
	8.1.3.3 D() [1/2]	64
	8.1.3.4 D() [2/2]	64
	8.1.3.5 D_ptr() [1/2]	64
	8.1.3.6 D_ptr() [2/2]	64
	8.1.3.7 default_val()	64
	8.1.3.8 flush_data()	64
	8.1.3.9 get_cell()	65
	8.1.3.10 get_col_vec() [1/2]	65
	8.1.3.11 get_col_vec() [2/2]	65
	8.1.3.12 get_entries()	65
	8.1.3.13 get_row_vec() [1/2]	65
	<b>8.1.3.14 get_row_vec()</b> [2/2]	66
	8.1.3.15 insert_cell() [1/3]	66
	<b>8.1.3.16 insert_cell()</b> [2/3]	66
	<b>8.1.3.17 insert_cell()</b> [3/3]	66
	8.1.3.18 is_dense()	66
	8.1.3.19 is_empty()	67
	8.1.3.20 ncol()	67
	8.1.3.21 nnozero()	67
	8.1.3.22 nrow()	67
	8.1.3.23 operator()() [1/2]	67
	<b>8.1.3.24</b> operator()() [2/2]	67
	8.1.3.25 operator*=()	68
	8.1.3.26 operator+=() [1/3]	68
	<b>8.1.3.27</b> operator+=() [2/3]	68
	<b>8.1.3.28</b> operator+=() [3/3]	68
	<b>8.1.3.29</b> operator-=() [1/3]	68
	<b>8.1.3.30</b> operator-=() [2/3]	68
	<b>8.1.3.31 operator-=()</b> [3/3]	69
	8.1.3.32 operator/=()	69
	8.1.3.33 operator=() [1/2]	69
	8.1.3.34 operator=() [2/2]	69
	8.1.3.35 operator==()	69
	8.1.3.36 out_of_range()	69
	8.1.3.37 print()	70
	8.1.3.38 print_n()	70
	8.1.3.39 reserve()	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
8.4.2 Constructor & Destructor Documentation
8.4.2.1 BArrayDense() [1/6]
8.4.2.2 BArrayDense() [2/6]
8.4.2.3 BArrayDense() [3/6]
8.4.2.4 BArrayDense() [4/6]
8.4.2.5 BArrayDense() [5/6]
8.4.2.6 BArrayDense() [6/6]
8.4.2.7 ~BArrayDense()
8.4.3 Member Function Documentation
8.4.3.1 clear()
8.4.3.2 col() [1/2] 84
8.4.3.3 col() [2/2]
8.4.3.4 colsum()
8.4.3.5 D() [1/2]
8.4.3.6 D() [2/2]
8.4.3.7 D_ptr() [1/2]
8.4.3.8 D_ptr() [2/2]
8.4.3.9 default_val()
8.4.3.10 get_cell()
8.4.3.11 get_col_vec() [1/2]
8.4.3.12 get_col_vec() [2/2]
8.4.3.13 get_data()
8.4.3.14 get_entries()
8.4.3.15 get_row_vec() [1/2]
8.4.3.16 get_row_vec() [2/2]
8.4.3.17 insert_cell() [1/2]
8.4.3.18 insert_cell() [2/2]
8.4.3.19 is_dense()
8.4.3.20 is_empty()
8.4.3.21 ncol()
8.4.3.22 nnozero()
8.4.3.23 nrow()
8.4.3.24 operator()() [1/2]
8.4.3.25 operator()() [2/2]
8.4.3.26 operator*=()
8.4.3.27 operator+=() [1/3]
8.4.3.28 operator+=() [2/3]
8.4.3.29 operator+=() [3/3] 90
8.4.3.30 operator-=() [1/3]
8.4.3.31 operator-=() [2/3]
8.4.3.32 operator-=() [3/3]

8.4.3.33 operator/=()	90
8.4.3.34 operator=() [1/2]	91
8.4.3.35 operator=() [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
8.4.3.42 row() [1/2]	92
8.4.3.43 row() [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	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 $\sim$ 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
8.15.3.2 add() [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 ConstBArrayRowlter< 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 ConstBArrayRowIter()	. 127
8.17.2.2 ~ConstBArrayRowlter()	. 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
<b>8.19.2.1 Counters()</b> [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
8.19.3.1 add_counter() [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
8.19.3.8 operator=() [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()
8.20.3.9 get_X()
8.20.3.10 get_X_names()
8.20.3.11 get_Y()
8.20.3.12 get_Y_names()
8.20.3.13 init()
8.20.3.14 is_motif()
8.20.3.15 logodds()
8.20.3.16 motif_census()
8.20.3.17 print()
8.20.3.18 set_names()
8.20.3.19 simulate()
8.21 DEFMCounterData Class Reference
8.21.1 Detailed Description
8.22 DEFMData Class Reference
8.22.1 Detailed Description
8.23 DEFMRuleData Class Reference
8.23.1 Detailed Description
8.24 DEFMRuleDynData Class Reference
8.24.1 Detailed Description
8.25 Entries < Cell_Type > Class Template Reference
8.25.1 Detailed Description
8.25.2 Constructor & Destructor Documentation
8.25.2.1 Entries() [1/2]
8.25.2.2 Entries() [2/2]
8.25.2.3 ~Entries()
8.25.3 Member Function Documentation
8.25.3.1 resize()
8.25.4 Member Data Documentation
8.25.4.1 source
8.25.4.2 target
8.25.4.3 val
8.26 Flock Class Reference
8.26.1 Detailed Description
8.26.2 Constructor & Destructor Documentation
8.26.2.1 Flock()
8.26.2.2 ~Flock()
8.26.3 Member Function Documentation
8.26.3.1 add_data()
8.26.3.2 colnames()
8.26.3.3 get_counters()
8.26.3.4 get_model()

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]	34
8.28.2.5 ~Geese()	34
8.28.3 Member Function Documentation	34
8.28.3.1 calc_reduced_sequence()	35
8.28.3.2 calc_sequence()	35
8.28.3.3 colnames()	35
8.28.3.4 get_annotated_nodes()	35
8.28.3.5 get_annotations()	35
8.28.3.6 get_counters()	36
8.28.3.7 get_model()	36
8.28.3.8 get_probabilities()	36
8.28.3.9 get_rengine()	36
8.28.3.10 get_states()	36
8.28.3.11 get_support_fun()	37
8.28.3.12 inherit_support()	37
8.28.3.13 init()	37
8.28.3.14 init_node()	37
8.28.3.15 likelihood()	37
8.28.3.16 likelihood_exhaust()	38
8.28.3.17 nannotations()	36
8.28.3.18 nfuns()	36
8.28.3.19 nleafs()	36
8.28.3.20 nnodes()	36
8.28.3.21 nterms()	39
8.28.3.22 observed_counts()	39
8.28.3.23 operator=() [1/2]	39
8.28.3.24 operator=() [2/2]	39
8.28.3.25 parse_polytomies()	39
8.28.3.26 predict()	70
8.28.3.27 predict_backend()	70
8.28.3.28 predict_exhaust()	70
8.28.3.29 predict_exhaust_backend()	70
8.28.3.30 predict_sim()	70
8.28.3.31 print()	71
8.28.3.32 print_nodes()	71
8.28.3.33 print_observed_counts()	71
8.28.3.34 set_seed()	71
8.28.3.35 simulate()	71
8.28.3.36 support_size()	71
8.28.3.37 update_annotations()	72
8.28.4 Member Data Documentation	72
8 28 / 1 delete rengine	72

	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
	Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class Tem-	
•	eference	
	Detailed Description	
	Constructor & Destructor Documentation	
	8.29.2.1 Model() [1/3]	
	8.29.2.2 Model() [2/3]	
	8.29.2.3 Model() [3/3]	
	8.29.2.4 ~Model()	
8.29.3 N	Member Function Documentation	
	8.29.3.1 add_array()	
	8.29.3.2 add_counter() [1/2]	
	<b>8.29.3.3</b> add_counter() [2/2]	181
	8.29.3.4 add_hasher()	181
	8.29.3.5 add_rule() [1/2]	181
	<b>8.29.3.6</b> add_rule() [2/2]	181
	<b>8.29.3.7</b> add_rule_dyn() [1/2]	182
	<b>8.29.3.8</b> add_rule_dyn() [2/2]	182
	8.29.3.9 colnames()	182
	8.29.3.10 conditional_prob()	182
	8.29.3.11 gen_key()	183
	8.29.3.12 get_arrays2support()	183
	8.29.3.13 get_counters()	183
	8.29.3.14 get_norm_const()	183
	8.29.3.15 get_pset()	184
	8.29.3.16 get_pset_arrays()	184
	8.29.3.17 get_pset_probs()	184
	8.29.3.18 get_pset_stats() [1/2]	184
	8.29.3.19 get_pset_stats() [2/2]	184
	8.29.3.20 get_rengine()	185
	8.29.3.21 get rules()	185

	3.29.3.22 get_rules_dyn()	5
	3.29.3.23 get_stats_support()	5
	3.29.3.24 get_stats_target()	5
	3.29.3.25 get_support_fun()	6
	3.29.3.26 likelihood() [1/4]	6
	3.29.3.27 likelihood() [2/4]	6
	3.29.3.28 likelihood() [3/4]	6
	3.29.3.29 likelihood() [4/4]	17
	3.29.3.30 likelihood_total()	37
	3.29.3.31 nrules()	17
	3.29.3.32 nrules_dyn()	37
	3.29.3.33 nterms()	17
	3.29.3.34 operator=()	8
	3.29.3.35 print()	8
	3.29.3.36 print_stats()	8
	3.29.3.37 sample() [1/2]	8
	3.29.3.38 sample() [2/2]	9
	3.29.3.39 set_counters()	9
	3.29.3.40 set_rengine()	9
	3.29.3.41 set_rules()	9
	3.29.3.42 set_rules_dyn()	9
	3.29.3.43 set_seed()	0
	3.29.3.44 set_transform_model()	0
	3.29.3.45 size()	0
	3.29.3.46 size_unique()	0
	3.29.3.47 store_psets()	1
	3.29.3.48 support_size()	1
	3.29.3.49 transform_model()	1
8.29.4 N	ember Data Documentation	1
	3.29.4.1 arrays2support	11
	3.29.4.2 counter_fun	11
	3.29.4.3 counters	12
	3.29.4.4 delete_counters	12
	3.29.4.5 delete_rengine	12
	3.29.4.6 delete_rules	12
	3.29.4.7 delete_rules_dyn	12
	3.29.4.8 first_calc_done	13
	3.29.4.9 keys2support	13
	3.29.4.10 normalizing_constants	13
	3.29.4.11 params_last	13
	3.29.4.12 pset_arrays	14
	3.29.4.13 pset_probs	)4

8.29.4.14 pset_stats	
8.29.4.15 rengine	
8.29.4.16 rules	
8.29.4.17 rules_dyn	
8.29.4.18 stats_support	195
8.29.4.19 stats_support_n_arrays	
8.29.4.20 stats_target	196
8.29.4.21 support_fun	196
8.29.4.22 transform_model_fun	196
8.29.4.23 transform_model_term_names	197
8.29.4.24 with_pset	197
8.30 NetCounterData Class Reference	197
8.30.1 Detailed Description	197
8.30.2 Constructor & Destructor Documentation	198
8.30.2.1 NetCounterData() [1/2]	198
8.30.2.2 NetCounterData() [2/2]	198
8.30.2.3 ~NetCounterData()	198
8.30.3 Member Data Documentation	198
8.30.3.1 indices	198
8.30.3.2 numbers	198
8.31 NetworkData Class Reference	199
8.31.1 Detailed Description	199
8.31.2 Constructor & Destructor Documentation	199
8.31.2.1 NetworkData() [1/3]	199
<b>8.31.2.2</b> NetworkData() [2/3]	199
<b>8.31.2.3</b> NetworkData() [3/3]	200
8.31.2.4 ~NetworkData()	200
8.31.3 Member Data Documentation	200
8.31.3.1 directed	200
8.31.3.2 vertex_attr	201
8.32 Node Class Reference	201
8.32.1 Detailed Description	202
8.32.2 Constructor & Destructor Documentation	202
<b>8.32.2.1 Node()</b> [1/5]	202
8.32.2.2 Node() [2/5]	
8.32.2.3 Node() [3/5]	
8.32.2.4 Node() [4/5]	
8.32.2.5 Node() [5/5]	
8.32.2.6 ~Node()	
8.32.3 Member Function Documentation	
8.32.3.1 get_parent()	
8.32.3.2 is leaf()	

8.32.3.3 noffspring()	04
8.32.4 Member Data Documentation	04
8.32.4.1 annotations	04
8.32.4.2 array	04
8.32.4.3 arrays	05
8.32.4.4 duplication	05
8.32.4.5 id	05
8.32.4.6 narray	05
8.32.4.7 offspring	05
8.32.4.8 ord	06
8.32.4.9 parent	06
8.32.4.10 probability	06
8.32.4.11 subtree_prob	06
8.32.4.12 visited	06
8.33 NodeData Class Reference	07
8.33.1 Detailed Description	07
8.33.2 Constructor & Destructor Documentation	07
8.33.2.1 NodeData()	07
8.33.3 Member Data Documentation	07
8.33.3.1 blengths	80
8.33.3.2 duplication	80
8.33.3.3 states	80
8.34 PhyloCounterData Class Reference	80
8.34.1 Detailed Description	09
8.34.2 Constructor & Destructor Documentation	09
8.34.2.1 PhyloCounterData() [1/2]	09
8.34.2.2 PhyloCounterData() [2/2]	09
8.34.3 Member Function Documentation	09
8.34.3.1 at()	09
8.34.3.2 begin()	09
8.34.3.3 empty()	10
8.34.3.4 end()	10
8.34.3.5 get_counters()	10
8.34.3.6 operator()()	10
8.34.3.7 operator[]()	10
8.34.3.8 push_back()	10
8.34.3.9 reserve()	11
8.34.3.10 shrink_to_fit()	11
8.34.3.11 size()	11
8.35 PhyloRuleDynData Class Reference	11
8.35.1 Detailed Description	12
8.35.2 Constructor & Destructor Documentation	12

8.35.2.1 PhyloRuleDynData()	12
8.35.2.2 ~PhyloRuleDynData()	12
8.35.3 Member Function Documentation	12
8.35.3.1 operator()()	12
8.35.4 Member Data Documentation	12
8.35.4.1 counts	12
8.35.4.2 duplication	13
8.35.4.3 lb	13
8.35.4.4 pos	13
8.35.4.5 ub	13
8.36 PowerSet< Array_Type, Data_Rule_Type > Class Template Reference	13
8.36.1 Detailed Description	14
8.36.2 Constructor & Destructor Documentation	15
8.36.2.1 PowerSet() [1/3]	15
8.36.2.2 PowerSet() [2/3]	15
8.36.2.3 PowerSet() [3/3]	15
8.36.2.4 ~PowerSet()	15
8.36.3 Member Function Documentation	15
8.36.3.1 add_rule() [1/2]	16
8.36.3.2 add_rule() [2/2] 2	16
8.36.3.3 begin()	16
8.36.3.4 calc()	16
8.36.3.5 end()	16
8.36.3.6 get_data()	17
8.36.3.7 get_data_ptr()	17
8.36.3.8 init_support()	17
8.36.3.9 operator[]()	17
8.36.3.10 reset()	17
8.36.3.11 size()	18
8.36.4 Member Data Documentation	18
8.36.4.1 coordinates_free	18
8.36.4.2 coordinates_locked	18
8.36.4.3 data	18
8.36.4.4 EmptyArray	18
8.36.4.5 M	19
8.36.4.6 N	19
8.36.4.7 n_free	19
8.36.4.8 n_locked	19
8.36.4.9 rules	19
8.36.4.10 rules_deleted	20
8.37 Progress Class Reference	20
8 37 1 Detailed Description	าวก

8.37.2 Constructor & Destructor Documentation	20
8.37.2.1 Progress()	20
8.37.2.2 ~Progress()	21
8.37.3 Member Function Documentation	21
8.37.3.1 end()	21
8.37.3.2 next()	21
8.38 Rule < Array_Type, Data_Type > Class Template Reference	21
8.38.1 Detailed Description	22
8.38.2 Constructor & Destructor Documentation	22
8.38.2.1 Rule() [1/2]	22
8.38.2.2 Rule() [2/2]	22
8.38.2.3 ~Rule()	23
8.38.3 Member Function Documentation	23
8.38.3.1 D()	23
8.38.3.2 get_description() [1/2]	23
8.38.3.3 get_description() [2/2]	23
8.38.3.4 get_name() [1/2]	23
8.38.3.5 get_name() [2/2]	24
8.38.3.6 operator()()	24
8.39 Rules < Array_Type, Data_Type > Class Template Reference	24
8.39.1 Detailed Description	25
8.39.2 Constructor & Destructor Documentation	25
8.39.2.1 Rules() [1/2]	25
8.39.2.2 Rules() [2/2]	25
8.39.2.3 ∼Rules()	26
8.39.3 Member Function Documentation	26
8.39.3.1 add_rule() [1/2]	26
8.39.3.2 add_rule() [2/2]	26
8.39.3.3 begin()	26
8.39.3.4 end()	26
8.39.3.5 get_descriptions()	27
8.39.3.6 get_names()	27
8.39.3.7 get_seq()	27
8.39.3.8 operator()()	27
8.39.3.9 operator=()	28
8.39.3.10 size()	28
8.40 StatsCounter< Array_Type, Data_Type > Class Template Reference	28
8.40.1 Detailed Description	
8.40.2 Constructor & Destructor Documentation	
8.40.2.1 StatsCounter() [1/3]	29
8.40.2.2 StatsCounter() [2/3]	30
8.40.2.3 StatsCounter() [3/3]	30

8.40.2.4 ~StatsCounter()	23
8.40.3 Member Function Documentation	23
8.40.3.1 add_counter()	230
8.40.3.2 count_all()	23
8.40.3.3 count_current()	23
8.40.3.4 count_init()	23
8.40.3.5 get_counters()	23
8.40.3.6 get_descriptions()	23
8.40.3.7 get_names()	23
8.40.3.8 reset_array()	23
8.40.3.9 set_counters()	23
8.40.3.10 size()	23
8.41 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class Te	
plate Reference	
8.41.1 Detailed Description	
8.41.2 Constructor & Destructor Documentation	
<b>8.41.2.1 Support()</b> [1/3]	
<b>8.41.2.2 Support()</b> [2/3]	23
<b>8.41.2.3 Support()</b> [3/3]	23
8.41.2.4 ~Support()	23
8.41.3 Member Function Documentation	23
8.41.3.1 add_counter()	23
<b>8.41.3.2</b> add_rule() [1/2]	23
<b>8.41.3.3 add_rule()</b> [2/2]	230
8.41.3.4 add_rule_dyn() [1/2]	230
<b>8.41.3.5 add_rule_dyn()</b> [2/2]	230
8.41.3.6 calc()	23
8.41.3.7 eval_rules_dyn()	23
8.41.3.8 get_counters()	23
8.41.3.9 get_counts()	23
8.41.3.10 get_current_stats()	238
8.41.3.11 get_data()	23
8.41.3.12 get_rules()	23
8.41.3.13 get_rules_dyn()	23
8.41.3.14 init_support()	239
8.41.3.15 print()	239
8.41.3.16 reset_array() [1/2]	239
<b>8.41.3.17 reset_array()</b> [2/2]	239
8.41.3.18 set_counters()	
8.41.3.19 set_rules()	
8.41.3.20 set_rules_dyn()	
8.41.4 Member Data Decumentation	24

8.41.4.1 change_stats	240
8.41.4.2 coordiantes_n_free	241
8.41.4.3 coordiantes_n_locked	241
8.41.4.4 coordinates_free	241
8.41.4.5 coordinates_locked	241
8.41.4.6 current_stats	241
8.41.4.7 delete_counters	242
8.41.4.8 delete_rules	242
8.41.4.9 delete_rules_dyn	242
8.41.4.10 hashes	242
8.41.4.11 hashes_initialized	242
8.41.4.12 M	243
8.41.4.13 max_num_elements	243
8.41.4.14 N	243
8.41.4.15 n_counters	243
8.42 vecHasher < T > Struct Template Reference	243
8.42.1 Detailed Description	244
8.42.2 Member Function Documentation	244
8.42.2.1 operator()()	244
9 File Documentation	245
9.1 include/parry/parray-pones.ripp file Reference	. 245
9.1 include/barry/barray-bones.hpp File Reference	
9.2 include/barry/barray-iterator.hpp File Reference	245
9.2 include/barry/barray-iterator.hpp File Reference 9.3 include/barry/barray-meat-operators.hpp File Reference	245
9.2 include/barry/barray-iterator.hpp File Reference 9.3 include/barry/barray-meat-operators.hpp File Reference 9.3.1 Macro Definition Documentation	245 246 246
9.2 include/barry/barray-iterator.hpp File Reference 9.3 include/barry/barray-meat-operators.hpp File Reference 9.3.1 Macro Definition Documentation 9.3.1.1 BARRAY_TEMPLATE	245 246 246 247
9.2 include/barry/barray-iterator.hpp File Reference 9.3 include/barry/barray-meat-operators.hpp File Reference 9.3.1 Macro Definition Documentation 9.3.1.1 BARRAY_TEMPLATE 9.3.1.2 BARRAY_TEMPLATE_ARGS	245 246 246 247 247
9.2 include/barry/barray-iterator.hpp File Reference 9.3 include/barry/barray-meat-operators.hpp File Reference 9.3.1 Macro Definition Documentation 9.3.1.1 BARRAY_TEMPLATE 9.3.1.2 BARRAY_TEMPLATE_ARGS 9.3.1.3 BARRAY_TYPE	245 246 246 247 247 247
9.2 include/barry/barray-iterator.hpp File Reference 9.3 include/barry/barray-meat-operators.hpp File Reference 9.3.1 Macro Definition Documentation 9.3.1.1 BARRAY_TEMPLATE 9.3.1.2 BARRAY_TEMPLATE_ARGS	245 246 246 247 247 247 247
9.2 include/barry/barray-iterator.hpp File Reference 9.3 include/barry/barray-meat-operators.hpp File Reference 9.3.1 Macro Definition Documentation 9.3.1.1 BARRAY_TEMPLATE 9.3.1.2 BARRAY_TEMPLATE_ARGS 9.3.1.3 BARRAY_TYPE 9.3.1.4 COL 9.3.1.5 ROW	245 246 246 247 247 247 247
9.2 include/barry/barray-iterator.hpp File Reference 9.3 include/barry/barray-meat-operators.hpp File Reference 9.3.1 Macro Definition Documentation 9.3.1.1 BARRAY_TEMPLATE 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	245 246 247 247 247 247 247 247
9.2 include/barry/barray-iterator.hpp File Reference 9.3 include/barry/barray-meat-operators.hpp File Reference 9.3.1 Macro Definition Documentation 9.3.1.1 BARRAY_TEMPLATE 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]	245 246 247 247 247 247 247 247 247 248
9.2 include/barry/barray-iterator.hpp File Reference 9.3 include/barry/barray-meat-operators.hpp File Reference 9.3.1 Macro Definition Documentation 9.3.1.1 BARRAY_TEMPLATE 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]	245 246 247 247 247 247 247 247 248 248
9.2 include/barry/barray-iterator.hpp File Reference 9.3 include/barry/barray-meat-operators.hpp File Reference 9.3.1 Macro Definition Documentation 9.3.1.1 BARRAY_TEMPLATE 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() [2/6]	245 246 247 247 247 247 247 247 248 248
9.2 include/barry/barray-iterator.hpp File Reference 9.3 include/barry/barray-meat-operators.hpp File Reference 9.3.1 Macro Definition Documentation 9.3.1.1 BARRAY_TEMPLATE 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] 9.3.2.3 BARRAY_TEMPLATE() [3/6]	245 246 247 247 247 247 247 247 248 248 248
9.2 include/barry/barray-iterator.hpp File Reference 9.3 include/barry/barray-meat-operators.hpp File Reference 9.3.1 Macro Definition Documentation 9.3.1.1 BARRAY_TEMPLATE 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() [4/6]	245 246 247 247 247 247 247 247 248 248 248 248
9.2 include/barry/barray-iterator.hpp File Reference 9.3 include/barry/barray-meat-operators.hpp File Reference 9.3.1 Macro Definition Documentation 9.3.1.1 BARRAY_TEMPLATE 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() [3/6] 9.3.2.5 BARRAY_TEMPLATE() [5/6] 9.3.2.5 BARRAY_TEMPLATE() [5/6]	245 246 247 247 247 247 247 247 248 248 248 248 248 248
9.2 include/barry/barray-iterator.hpp File Reference 9.3 include/barry/barray-meat-operators.hpp File Reference 9.3.1 Macro Definition Documentation 9.3.1.1 BARRAY_TEMPLATE 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() [6/6]	245 246 247 247 247 247 247 248 248 248 248 248 248 249
9.2 include/barry/barray-iterator.hpp File Reference 9.3 include/barry/barray-meat-operators.hpp File Reference 9.3.1 Macro Definition Documentation 9.3.1.1 BARRAY_TEMPLATE 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() [3/6] 9.3.2.5 BARRAY_TEMPLATE() [4/6] 9.3.2.6 BARRAY_TEMPLATE() [6/6] 9.3.2.7 BARRAY_TEMPLATE() [6/6]	245 246 247 247 247 247 247 248 248 248 248 248 249 249
9.2 include/barry/barray-iterator.hpp File Reference 9.3 include/barry/barray-meat-operators.hpp File Reference 9.3.1 Macro Definition Documentation 9.3.1.1 BARRAY_TEMPLATE 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() [5/6] 9.3.2.7 BARRAY_TEMPLATE() [6/6]	245 246 247 247 247 247 247 248 248 248 248 249 249 249

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/barry/barraydense-meat.hpp 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]
<b>9.15.2.3 BROW_TEMPLATE()</b> [3/5]
9.15.2.4 BROW_TEMPLATE() [4/5]
<b>9.15.2.5 BROW_TEMPLATE()</b> [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_ONE
9.20.1.2 BARRY_ONE_DENSE
9.20.1.3 BARRY_UNUSED
9.20.1.4 BARRY_ZERO
9.20.1.5 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]
<b>9.26.2.4 COUNTER_TEMPLATE()</b> [3/9]
9.26.2.5 COUNTER_TEMPLATE() [4/9]
<b>9.26.2.6 COUNTER_TEMPLATE()</b> [5/9]
9.26.2.7 COUNTER_TEMPLATE() [6/9]
9.26.2.8 COUNTER_TEMPLATE() [7/9]
9.26.2.9 COUNTER_TEMPLATE() [8/9]
9.26.2.10 COUNTER_TEMPLATE() [9/9]
9.26.2.11 COUNTERS_TEMPLATE() [1/9]
9.26.2.12 COUNTERS_TEMPLATE() [2/9]
9.26.2.13 COUNTERS_TEMPLATE() [3/9]
9.26.2.14 COUNTERS_TEMPLATE() [4/9]
9.26.2.15 COUNTERS_TEMPLATE() [5/9]
9.26.2.16 COUNTERS_TEMPLATE() [6/9]
9.26.2.17 COUNTERS_TEMPLATE() [7/9]
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()

9.26.2.22 for()	279
9.26.2.23 hasher() [1/2]	280
9.26.2.24 hasher() [2/2]	280
9.26.2.25 hasher_fun() [1/2]	280
9.26.2.26 hasher_fun() [2/2]	280
9.26.2.27 if() [1/3]	280
9.26.2.28 if() [2/3]	
9.26.2.29 if() [3/3]	
9.26.2.30 init_fun() [1/3]	281
9.26.2.31 init_fun() [2/3]	281
9.26.2.32 init_fun() [3/3]	
9.26.2.33 name()	
9.26.3 Variable Documentation	281
9.26.3.1 add_dims	281
9.26.3.2 count_fun	282
9.26.3.3 counter	282
9.26.3.4 counter	282
9.26.3.5 data	282
9.26.3.6 desc	283
9.26.3.7 fun	283
9.26.3.8 fun	283
9.26.3.9 hasher_fun	283
9.26.3.10 i	284
9.26.3.11 init_fun	284
9.26.3.12 j	284
9.26.3.13 name	284
9.26.3.14 noexcept	284
9.26.3.15 res	285
9.26.3.16 return	285
9.27 include/barry/counters/network-css.hpp File Reference	285
9.27.1 Macro Definition Documentation	286
9.27.1.1 CSS_APPEND	287
9.27.1.2 CSS_CASE_ELSE	287
9.27.1.3 CSS_CASE_PERCEIVED	287
9.27.1.4 CSS_CASE_TRUTH	287
9.27.1.5 CSS_CHECK_SIZE	287
9.27.1.6 CSS_CHECK_SIZE_INIT	288
9.27.1.7 CSS_MATCH_TYPE	288
9.27.1.8 CSS_NET_COUNTER_LAMBDA_INIT	288
9.27.1.9 CSS_PERCEIVED_CELLS	288
9.27.1.10 CSS_SIZE	289
9.27.1.11 CSS TRUE CELLS	289

9.27.2 Function Documentation
9.27.2.1 counter_css_census01()
9.27.2.2 counter_css_census02()
9.27.2.3 counter_css_census03()
9.27.2.4 counter_css_census04()
9.27.2.5 counter_css_census05()
9.27.2.6 counter_css_census06()
9.27.2.7 counter_css_census07()
9.27.2.8 counter_css_census08()
9.27.2.9 counter_css_census09()
9.27.2.10 counter_css_census10()
9.27.2.11 counter_css_completely_false_recip_comiss()
9.27.2.12 counter_css_completely_false_recip_omiss()
9.27.2.13 counter_css_mixed_recip()
9.27.2.14 counter_css_partially_false_recip_commi()
9.27.2.15 counter_css_partially_false_recip_omiss()
9.28 include/barry/counters/network.hpp File Reference
9.28.1 Macro Definition Documentation
9.28.1.1 BARRY_ZERO_NETWORK
9.28.1.2 BARRY_ZERO_NETWORK_DENSE
9.28.1.3 NET_C_DATA_IDX
9.28.1.4 NET_C_DATA_NUM
9.28.1.5 NETWORK_COUNTER
9.28.1.6 NETWORK_COUNTER_LAMBDA
9.28.1.7 NETWORK_RULE
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 Macro Definition Documentation
9.31.1.1 MODEL_TEMPLATE
9.31.1.2 MODEL_TEMPLATE_ARGS
9.31.1.3 MODEL_TYPE
9.31.2 Function Documentation
9.31.2.1 for() [1/2]
9.31.2.2 for() [2/2]
9.31.2.3 if() [1/4]
9.31.2.4 if() [2/4]
9.31.2.5 if() [3/4]
9.31.2.6 if() [4/4]
9.31.2.7 insert_cell()
9.31.2.8 likelihood_()
9.31.2.9 MODEL_TEMPLATE() [1/33]
9.31.2.10 MODEL_TEMPLATE() [2/33]
<b>9.31.2.11 MODEL_TEMPLATE()</b> [3/33]
9.31.2.12 MODEL_TEMPLATE() [4/33]
<b>9.31.2.13 MODEL_TEMPLATE()</b> [5/33]
<b>9.31.2.14 MODEL_TEMPLATE()</b> [6/33]
<b>9.31.2.15 MODEL_TEMPLATE()</b> [7/33]
<b>9.31.2.16 MODEL_TEMPLATE()</b> [8/33]
<b>9.31.2.17 MODEL_TEMPLATE()</b> [9/33]
9.31.2.18 MODEL_TEMPLATE() [10/33]
9.31.2.19 MODEL_TEMPLATE() [11/33]
9.31.2.20 MODEL_TEMPLATE() [12/33]
9.31.2.21 MODEL_TEMPLATE() [13/33]
9.31.2.22 MODEL_TEMPLATE() [14/33]
9.31.2.23 MODEL_TEMPLATE() [15/33]
9.31.2.24 MODEL_TEMPLATE() [16/33]
9.31.2.25 MODEL_TEMPLATE() [17/33]
9.31.2.26 MODEL_TEMPLATE() [18/33]
9.31.2.27 MODEL_TEMPLATE() [19/33]
9.31.2.28 MODEL_TEMPLATE() [20/33]
9.31.2.29 MODEL_TEMPLATE() [21/33]
9.31.2.30 MODEL_TEMPLATE() [22/33]
9.31.2.31 MODEL_TEMPLATE() [23/33]
9.31.2.32 MODEL_TEMPLATE() [24/33]
<b>9.31.2.33 MODEL_TEMPLATE()</b> [25/33]
9.31.2.34 MODEL_TEMPLATE() [26/33]
9.31.2.35 MODEL_TEMPLATE() [27/33]
<b>9.31.2.36 MODEL_TEMPLATE()</b> [28/33]
9.31.2.37 MODEL_TEMPLATE() [29/33]

9	31.2.38 MODEL_TEMPLATE() [30/33]	11
9	31.2.39 MODEL_TEMPLATE() [31/33]	12
9	31.2.40 MODEL_TEMPLATE() [32/33]	12
9	31.2.41 MODEL_TEMPLATE() [33/33]	12
9	31.2.42 push_back() [1/2]	12
9	31.2.43 push_back() [2/2]	12
9	31.2.44 reserve() [1/2]	12
9	31.2.45 reserve() [2/2]	13
9	31.2.46 return()	13
9	31.2.47 set_counters()	13
9	31.2.48 set_rules()	13
9	31.2.49 set_rules_dyn()	13
9	31.2.50 size()	13
9	31.2.51 update_normalizing_constant() 31	14
9	31.2.52 urand()	14
9.31.3 Va	iable Documentation	14
9	31.3.1 a	14
9	31.3.2 count_fun	14
9	31.3.3 counter	14
9	31.3.4 counters	15
9	31.3.5 cumprob	15
9	31.3.6 data	15
9	31.3.7 Data_Counter_Type	15
9	31.3.8 Data_Rule_Type	15
9	31.3.9 delete_rules	16
9	31.3.10 delete_rules_dyn	16
9	31.3.11 else	16
9	31.3.12 force_new	16
9	31.3.13 fun	16
9	31.3.14 i	17
9	31.3.15 i_matches	17
9	31.3.16 init_fun	17
9	31.3.17 j	17
9	31.3.18 k	17
9	31.3.19 key	17
9	31.3.20 locator	18
9	31.3.21 params	18
9	31.3.22 probs	18
9	31.3.23 pset_arrays	18
9	31.3.24 r	18
9	31.3.25 return	18
9	31.3.26 rule_fun	19

0.21.2.27 rules
9.31.3.27 rules
9.31.3.29 rules_dyn
9.31.3.30 stats
9.31.3.31 stats_support_n_arrays
- +
9.31.3.32 temp_stats
9.31.3.33 tmp_counts
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()
9.39 include/barry/models/geese.hpp File Reference
9.40 include/barry/models/geese/flock-bones.hpp File Reference
9.41 include/barry/models/geese/flock-meat.hpp File Reference
9.42 include/barry/models/geese/geese-bones.hpp File Reference
9.42.1 Macro Definition Documentation
9.42.1.1 INITIALIZED
9.42.2 Function Documentation
9.42.2.1 keygen_full()
9.42.2.2 RULE_FUNCTION()
9.42.2.3 vec_diff()
9.42.2.4 vector_caster()
9.42.2.4 vector_caster()
-a 4a include/batry/models/deese/deese-meat-constructors flob file Belefence 334

9.44 include/barry/models/geese/geese-meat-likelihood.hpp File Reference
9.45 include/barry/models/geese/geese-meat-likelihood_exhaust.hpp File Reference
9.46 include/barry/models/geese/geese-meat-predict.hpp File Reference
9.47 include/barry/models/geese/geese-meat-predict_exhaust.hpp File Reference
9.48 include/barry/models/geese/geese-meat-predict_sim.hpp File Reference
9.49 include/barry/models/geese/geese-meat-simulate.hpp File Reference
9.50 include/barry/models/geese/geese-meat.hpp File Reference
9.51 include/barry/models/geese/geese-node-bones.hpp File Reference
9.52 include/barry/models/geese/geese-types.hpp File Reference
9.52.1 Typedef Documentation
9.52.1.1 PhyloArray
9.52.1.2 PhyloCounter
9.52.1.3 PhyloCounters
9.52.1.4 PhyloModel
9.52.1.5 PhyloPowerSet
9.52.1.6 PhyloRule
9.52.1.7 PhyloRuleData
9.52.1.8 PhyloRuleDyn
9.52.1.9 PhyloRules
9.52.1.10 PhyloRulesDyn
9.52.1.11 PhyloStatsCounter
9.52.1.12 PhyloSupport
9.53 include/barry/powerset-bones.hpp File Reference
9.54 include/barry/powerset-meat.hpp File Reference
9.55 include/barry/progress.hpp File Reference
9.55.1 Macro Definition Documentation
9.55.1.1 BARRY_PROGRESS_BAR_WIDTH
9.56 include/barry/rules-bones.hpp File Reference
9.56.1 Function Documentation
9.56.1.1 rule_fun_default()
9.57 include/barry/rules-meat.hpp File Reference
9.58 include/barry/statscounter-bones.hpp File Reference
9.59 include/barry/statscounter-meat.hpp File Reference
9.59.1 Macro Definition Documentation
9.59.1.1 STATSCOUNTER_TEMPLATE
9.59.1.2 STATSCOUNTER_TEMPLATE_ARGS
9.59.1.3 STATSCOUNTER_TYPE
9.59.2 Function Documentation
9.59.2.1 clear()
9.59.2.2 for()
9.59.2.3 resize()
9.59.2.4 STATSCOUNTER_TEMPLATE() [1/9]

359

<b>9.59.2.5 STATSCOUNTER_TEMPLATE()</b> [2/9]	348
<b>9.59.2.6 STATSCOUNTER_TEMPLATE()</b> [3/9]	348
<b>9.59.2.7 STATSCOUNTER_TEMPLATE()</b> [4/9]	348
<b>9.59.2.8 STATSCOUNTER_TEMPLATE()</b> [5/9]	349
9.59.2.9 STATSCOUNTER_TEMPLATE() [6/9]	349
<b>9.59.2.10 STATSCOUNTER_TEMPLATE()</b> [7/9]	349
<b>9.59.2.11 STATSCOUNTER_TEMPLATE()</b> [8/9]	349
<b>9.59.2.12 STATSCOUNTER_TEMPLATE()</b> [9/9]	349
9.59.3 Variable Documentation	349
9.59.3.1 counter	350
9.59.3.2 counter_deleted	350
9.59.3.3 counters	350
9.59.3.4 counters	350
9.59.3.5 current_stats	350
9.59.3.6 EmptyArray	351
9.59.3.7 f	351
9.59.3.8 j	351
9.59.3.9 return	351
9.60 include/barry/support-bones.hpp File Reference	351
9.61 include/barry/support-meat.hpp File Reference	352
9.61.1 Macro Definition Documentation	352
9.61.1.1 BARRY_SUPPORT_MEAT_HPP	352
9.62 include/barry/typedefs.hpp File Reference	353
9.62.1 Typedef Documentation	355
9.62.1.1 Col_type	355
9.62.1.2 Counter_fun_type	355
9.62.1.3 Counts_type	355
9.62.1.4 Hasher_fun_type	355
9.62.1.5 MapVec_type	356
9.62.1.6 Row_type	356
9.62.1.7 Rule_fun_type	356
9.62.2 Function Documentation	356
9.62.2.1 sort_array()	356
9.62.2.2 vec_equal()	357
9.62.2.3 vec_equal_approx()	357
9.62.2.4 vec_inner_prod() [1/2]	358
9.62.2.5 vec_inner_prod() [2/2]	358
9.63 README.md File Reference	358

Index

# **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	197
NetworkData	
Data class for Networks	199
Node	
A single node for the model	201
NodeData	
Data definition for the PhyloArray class	207
PhyloCounterData	208
PhyloRuleDynData	211
PowerSet < Array_Type, Data_Rule_Type >	
Powerset of a binary array	213
Progress	
A simple progress bar	220
Rule < Array_Type, Data_Type >	
Rule for determining if a cell should be included in a sequence	221
Rules < Array_Type, Data_Type >	
Vector of objects of class Rule	224
StatsCounter< Array_Type, Data_Type >	
Count stats for a single Array	228
Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >	
Compute the support of sufficient statistics	232
vecHasher< T >	243

# **Chapter 5**

# File Index

# 5.1 File List

Here is a list of all files with brief descriptions:

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
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\_COUNTER\_LAMBDA(a)

Extension of a simple counter.

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

Overall functional gains.

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

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

k genes gain function nfun

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

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

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

Keeps track of how many pairs of genes preserve pseudostate.

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

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

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

Overall functional loss.

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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:

Value:

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

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

Definition at line 21 of file counters.hpp.

#### 6.1.2.7 MAKE\_DUPL\_VARS

```
#define MAKE_DUPL_VARS( )
```

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

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

Definition at line 12 of file counters.hpp.

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 42 of file counters.hpp.

# 6.1.2.9 PHYLO\_COUNTER\_LAMBDA

Extension of a simple counter.

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

Definition at line 36 of file counters.hpp.

#### 6.1.2.10 PHYLO RULE DYN LAMBDA

Definition at line 39 of file counters.hpp.

#### 6.1.3 Function Documentation

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

Function co-opting.

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

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

This algorithm implements the change statistic.

Definition at line 1296 of file counters.hpp.

#### 6.1.3.2 counter\_cogain()

Co-evolution (joint gain or loss)

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

Definition at line 791 of file counters.hpp.

#### 6.1.3.3 counter gains()

Functional gains for a specific function (nfun).

Definition at line 96 of file counters.hpp.

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 1630 of file counters.hpp.

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

k genes gain function nfun

Definition at line 156 of file counters.hpp.

# 6.1.3.6 counter\_genes\_changing()

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

Definition at line 228 of file counters.hpp.

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

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

Definition at line 1394 of file counters.hpp.

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

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

< How many genes diverge the parent

Definition at line 1514 of file counters.hpp.

#### 6.1.3.9 counter\_longest()

Longest branch mutates (either by gain or by loss)

Definition at line 848 of file counters.hpp.

# 6.1.3.10 counter\_loss()

Total count of losses for an specific function.

Definition at line 591 of file counters.hpp.

#### 6.1.3.11 counter\_maxfuns()

Cap the number of functions per gene.

Definition at line 529 of file counters.hpp.

6.1 Counting 21

#### 6.1.3.12 counter\_neofun()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1018 of file counters.hpp.

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

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1163 of file counters.hpp.

# 6.1.3.14 counter\_overall\_changes()

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

Definition at line 643 of file counters.hpp.

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

Overall functional gains.

Total number of gains (irrespective of the function).

Definition at line 58 of file counters.hpp.

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

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

Needs to specify function a.

Definition at line 1696 of file counters.hpp.

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

Overall functional loss.

Definition at line 481 of file counters.hpp.

#### 6.1.3.18 counter pairwise first gain()

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

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

Definition at line 1948 of file counters.hpp.

#### 6.1.3.19 counter pairwise neofun singlefun()

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

Definition at line 1099 of file counters.hpp.

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 1744 of file counters.hpp.

#### 6.1.3.21 counter pairwise preserving()

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

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

Definition at line 1809 of file counters.hpp.

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

Keeps track of how many pairs of genes preserve pseudostate.

Definition at line 297 of file counters.hpp.

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

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

Definition at line 379 of file counters.hpp.

#### 6.1.3.24 counter\_subfun()

Total count of Sub-functionalization events.

It requires to specify data = {funA, funB}

Definition at line 702 of file counters.hpp.

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

Definition at line 45 of file counters.hpp.

# 6.2 Statistical Models

Statistical models available in barry.

#### Classes

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

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

class Flock

A Flock is a group of Geese.

· class Geese

Annotated Phylo Model.

# 6.2.1 Detailed Description

Statistical models available in barry.

# 6.3 Network counters

Counters for network models.

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="">).</network,>
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\_dyn\_limit\_changes (PhyloSupport \*support, size\_t pos, size\_t lb, size\_t ub, size\_← t duplication=Geese::etype\_default)

Overall functional gains.

### **Variables**

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

Vector of covariates (complete vector)

size\_t DEFMData::obs\_start

Index of the observation in the data.

size\_t DEFMData::X\_ncol

Number of columns in the array of covariates.

size\_t DEFMData::X\_nrow

Number of rows in the array of covariates.

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

Value where the sorting of the covariates is stored.

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

6.4 Phylo rules 37

If false, then is a logit intercept.

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

# Convenient typedefs for network objects.

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

#### 6.4.1 Detailed Description

Rules for phylogenetic modeling.

#### **Parameters**

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

#### 6.4.2 Typedef Documentation

#### 6.4.2.1 DEFMCounter

typedef barry::Counter<DEFMArray, DEFMCounterData > DEFMCounter

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

# 6.4.2.2 DEFMCounters

typedef barry::Counters<DEFMArray, DEFMCounterData> DEFMCounters

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

#### 6.4.2.3 DEFMModel

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

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

#### 6.4.2.4 DEFMRule

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

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

#### 6.4.2.5 DEFMRuleDyn

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

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

#### 6.4.2.6 DEFMRules

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

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

#### 6.4.2.7 DEFMRulesDyn

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

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

#### 6.4.2.8 DEFMStatsCounter

typedef barry::StatsCounter<DEFMArray, DEFMCounterData> DEFMStatsCounter

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

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 2058 of file counters.hpp.

### 6.4.3.21 ∼DEFMCounterData()

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

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

#### 6.4.3.22 ∼DEFMData()

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

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

#### 6.4.3.23 ~DEFMRuleDynData()

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

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

# 6.4.4 Variable Documentation

#### 6.4.4.1 array

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

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

#### 6.4.4.2 counts

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

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

#### 6.4.4.3 covar\_sort

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

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

# 6.4.4.4 covar\_used

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

Value where the sorting of the covariates is stored.

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

#### 6.4.4.5 covariates

const double\* DEFMData::covariates

Vector of covariates (complete vector)

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

6.4 Phylo rules 45

#### 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

- 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)
- $\bullet \ \ void\ counter\_longest\ (PhyloCounters\ *counters,\ size\_t\ duplication=Geese::etype\_default)$ 
  - 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 1296 of file counters.hpp.

#### 6.5.2.2 counter\_cogain()

Co-evolution (joint gain or loss)

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

Definition at line 791 of file counters.hpp.

### 6.5.2.3 counter\_gains()

Functional gains for a specific function (nfun).

Definition at line 96 of file counters.hpp.

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

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

Needs to specify function a.

Definition at line 1630 of file counters.hpp.

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

k genes gain function nfun

Definition at line 156 of file counters.hpp.

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

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

Definition at line 228 of file counters.hpp.

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

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

Definition at line 1394 of file counters.hpp.

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

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

< How many genes diverge the parent

Definition at line 1514 of file counters.hpp.

6.5 Phylo counters 51

#### 6.5.2.9 counter\_longest()

Longest branch mutates (either by gain or by loss)

Definition at line 848 of file counters.hpp.

#### 6.5.2.10 counter\_loss()

Total count of losses for an specific function.

Definition at line 591 of file counters.hpp.

#### 6.5.2.11 counter\_maxfuns()

Cap the number of functions per gene.

Definition at line 529 of file counters.hpp.

# 6.5.2.12 counter\_neofun()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1018 of file counters.hpp.

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

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1163 of file counters.hpp.

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

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

Definition at line 643 of file counters.hpp.

# 6.5.2.15 counter\_overall\_gains()

Overall functional gains.

Total number of gains (irrespective of the function).

Definition at line 58 of file counters.hpp.

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

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

Needs to specify function a.

Definition at line 1696 of file counters.hpp.

6.5 Phylo counters 53

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

Overall functional loss.

Definition at line 481 of file counters.hpp.

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

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

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

Definition at line 1948 of file counters.hpp.

# 6.5.2.19 counter\_pairwise\_neofun\_singlefun()

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

Definition at line 1099 of file counters.hpp.

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

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

Needs to specify function a.

Definition at line 1744 of file counters.hpp.

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

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

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

Definition at line 1809 of file counters.hpp.

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

Keeps track of how many pairs of genes preserve pseudostate.

Definition at line 297 of file counters.hpp.

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

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

Definition at line 379 of file counters.hpp.

#### 6.5.2.24 counter\_subfun()

Total count of Sub-functionalization events.

It requires to specify data = {funA, funB}

Definition at line 702 of file counters.hpp.

# **Chapter 7**

# **Namespace Documentation**

# 7.1 barry Namespace Reference

barry: Your go-to motif accountant

# **Namespaces**

counters

Tree class and Treelterator class.

# 7.1.1 Detailed Description

barry: Your go-to motif accountant

# 7.2 barry::counters Namespace Reference

Tree class and Treelterator class.

# **Namespaces**

network

# 7.2.1 Detailed Description

Tree class and Treelterator class.

# 7.3 barry::counters::network Namespace Reference

# 7.4 CHECK Namespace Reference

Integer constants used to specify which cell should be check.

# **Variables**

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

Integer constants used to specify which cell should be check.

#### 7.4.2 Variable Documentation

#### 7.4.2.1 BOTH

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

Definition at line 27 of file typedefs.hpp.

# 7.4.2.2 NONE

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

Definition at line 28 of file typedefs.hpp.

#### 7.4.2.3 ONE

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

Definition at line 29 of file typedefs.hpp.

# 7.4.2.4 TWO

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

Definition at line 30 of file typedefs.hpp.

# 7.5 defm Namespace Reference

# 7.6 EXISTS Namespace Reference

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

#### **Variables**

```
• const int BOTH = -1
```

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

# 7.6.1 Detailed Description

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

# 7.6.2 Variable Documentation

# 7.6.2.1 AS\_ONE

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

Definition at line 45 of file typedefs.hpp.

# 7.6.2.2 AS\_ZERO

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

Definition at line 44 of file typedefs.hpp.

# 7.6.2.3 BOTH

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

Definition at line 38 of file typedefs.hpp.

#### 7.6.2.4 NONE

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

Definition at line 39 of file typedefs.hpp.

#### 7.6.2.5 ONE

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

Definition at line 40 of file typedefs.hpp.

# 7.6.2.6 TWO

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

Definition at line 41 of file typedefs.hpp.

#### 7.6.2.7 UKNOWN

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

Definition at line 43 of file typedefs.hpp.

# 7.7 geese Namespace Reference

# **Chapter 8**

# **Class Documentation**

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

Baseline class for binary arrays.

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

#### **Public Member Functions**

- bool operator== (const BArray< Cell\_Type, Data\_Type > &Array\_)
- ∼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()

#### 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	target When true tries to add repeated observations.	
target		
value		

• 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 94 of file barraydense-meat.hpp.

## 8.4.2.5 BArrayDense() [5/6]

Copy constructor.

Definition at line 150 of file barraydense-meat.hpp.

## 8.4.2.6 BArrayDense() [6/6]

Move operator.

Definition at line 244 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 325 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 920 of file barraydense-meat.hpp.

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

Definition at line 513 of file barraydense-meat.hpp.

## 8.4.3.3 col() [2/2]

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

# 8.4.3.4 colsum()

Definition at line 1023 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 360 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 365 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 350 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 355 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 586 of file barraydense-meat.hpp.

#### 8.4.3.10 get\_cell()

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

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

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

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

Definition at line 457 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 1013 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 525 of file barraydense-meat.hpp.

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

Definition at line 406 of file barraydense-meat.hpp.

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

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

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

Definition at line 712 of file barraydense-meat.hpp.

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

Definition at line 678 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 554 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 571 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 575 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 567 of file barraydense-meat.hpp.

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

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

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

Definition at line 641 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 591 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 610 of file barraydense-meat.hpp.

# 8.4.3.33 operator/=()

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

Move assignment.

Definition at line 262 of file barraydense-meat.hpp.

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

Assignment constructor.

Definition at line 194 of file barraydense-meat.hpp.

## 8.4.3.36 operator==()

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

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

Definition at line 370 of file barraydense-meat.hpp.

## 8.4.3.38 print()

Definition at line 979 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 970 of file barraydense-meat.hpp.

#### 8.4.3.40 resize()

Definition at line 934 of file barraydense-meat.hpp.

## 8.4.3.41 rm\_cell()

Definition at line 655 of file barraydense-meat.hpp.

# 8.4.3.42 row() [1/2]

Definition at line 485 of file barraydense-meat.hpp.

## 8.4.3.43 row() [2/2]

Definition at line 472 of file barraydense-meat.hpp.

#### 8.4.3.44 rowsum()

Definition at line 1018 of file barraydense-meat.hpp.

## 8.4.3.45 set\_data()

Set the data object.

#### **Parameters**

data_	
delete_←	
data_	

Definition at line 334 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 743 of file barraydense-meat.hpp.

## 8.4.3.47 swap\_cols()

Definition at line 827 of file barraydense-meat.hpp.

#### 8.4.3.48 swap\_rows()

Definition at line 799 of file barraydense-meat.hpp.

## 8.4.3.49 toggle\_cell()

Definition at line 780 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 892 of file barraydense-meat.hpp.

## 8.4.3.52 zero\_col()

Definition at line 873 of file barraydense-meat.hpp.

#### 8.4.3.53 zero\_row()

Definition at line 854 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()

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< std::vector< double > > \* get stats support ()
- std::vector< std::vector< double > > \* get\_stats\_target ()
- PhyloModel \* get\_model ()

Returns the joint likelihood of the model.

• Geese \* operator() (size\_t i, bool check\_bounds=true)

Access the i-th geese element.

### Information about the model

- size\_t nfuns () const noexcept
- size\_t ntrees () const noexcept
- std::vector< size\_t > nnodes () const noexcept
- std::vector< size\_t > nleafs () const noexcept
- size\_t nterms () const
- size\_t support\_size () const noexcept
- std::vector< std::string > colnames () const
- $\bullet \ \ \text{size\_t parse\_polytomies} \ (\text{bool verb=true}, \ \text{std::vector} < \ \text{size\_t} > * \ \text{dist=nullptr}) \ \text{const noexcept}$

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 224 of file flock-meat.hpp.

# 8.26.3.3 get\_counters()

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

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

# 8.26.3.4 get\_model()

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

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

# 8.26.3.5 get\_stats\_support()

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

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

8.26 Flock Class Reference 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 167 of file flock-meat.hpp.

### 8.26.3.11 nleafs()

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

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

### 8.26.3.12 nnodes()

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

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

# 8.26.3.13 nterms()

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

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

# 8.26.3.14 ntrees()

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

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

# 8.26.3.15 operator()()

Access the i-th geese element.

8.26 Flock Class Reference 155

### **Parameters**

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

Returns

Geese \*

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

# 8.26.3.16 parse\_polytomies()

Check polytomies and return the largest.

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

# 8.26.3.17 print()

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

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

# 8.26.3.18 set\_seed()

Set the seed of the model.

**Parameters** 

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

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

# 8.26.3.19 support\_size()

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

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

# 8.26.4 Member Data Documentation

# 8.26.4.1 dat

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

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

# 8.26.4.2 initialized

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

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

#### 8.26.4.3 model

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

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

# 8.26.4.4 nfunctions

```
size_t Flock::nfunctions = 0u
```

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

### 8.26.4.5 rengine

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

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

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

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

# 8.27 FreqTable < T > Class Template Reference

Frequency table of vectors.

```
#include <freqtable.hpp>
```

#### **Public Member Functions**

```
• FreqTable ()
```

- ∼FreqTable ()
- size\_t add (const std::vector< T > &x, size\_t \*h\_precomp)
- Counts\_type as\_vector () const
- const std::vector< double > & get\_data () const
- const std::unordered\_map< size\_t, size\_t > & get\_index () const
- void clear ()
- void reserve (size\_t n, size\_t k)
- void print () const
- size\_t size () const noexcept

Number of unique elements in the table. (.

size\_t make\_hash (const std::vector< T > &x) const

# 8.27.1 Detailed Description

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

Frequency table of vectors.

This is mostly used in Support. The main data is contained in the data double vector. The matrix is stored in a row-wise fashion, where the first element is the frequency with which the vector is observed.

For example, in a model with k terms the first k+1 elements of data would be:

- · weights
- term 1
- term 2
- ...
- · term k

Definition at line 22 of file freqtable.hpp.

# 8.27.2 Constructor & Destructor Documentation

# 8.27.2.1 FreqTable()

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

Definition at line 34 of file freqtable.hpp.

### 8.27.2.2 ∼FreqTable()

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

Definition at line 35 of file freqtable.hpp.

### 8.27.3 Member Function Documentation

# 8.27.3.1 add()

Definition at line 59 of file freqtable.hpp.

# 8.27.3.2 as\_vector()

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

Definition at line 139 of file freqtable.hpp.

### 8.27.3.3 clear()

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

Definition at line 168 of file freqtable.hpp.

### 8.27.3.4 get\_data()

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

Definition at line 40 of file freqtable.hpp.

# 8.27.3.5 get\_index()

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

Definition at line 41 of file freqtable.hpp.

### 8.27.3.6 make\_hash()

Definition at line 239 of file freqtable.hpp.

# 8.27.3.7 print()

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

Definition at line 204 of file freqtable.hpp.

### 8.27.3.8 reserve()

Definition at line 182 of file freqtable.hpp.

#### 8.27.3.9 size()

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

Number of unique elements in the table. (.

Returns

size\_t

Definition at line 231 of file freqtable.hpp.

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

• include/barry/freqtable.hpp

# 8.28 Geese Class Reference

Annotated Phylo Model.

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

# **Public Member Functions**

- ∼Geese ()
- void init (size\_t bar\_width=BARRY\_PROGRESS\_BAR\_WIDTH)
- void inherit\_support (const Geese &model\_, bool delete\_support\_=false)
- void calc\_sequence (Node \*n=nullptr)
- void calc\_reduced\_sequence ()
- double likelihood (const std::vector< double > &par, bool as\_log=false, bool use\_reduced\_sequence=true)
- double likelihood\_exhaust (const std::vector< double > &par)
- std::vector< double > get\_probabilities () const
- void set\_seed (const size\_t &s)
- std::vector< std::vector< size\_t >> simulate (const std::vector< double > &par)
- std::vector< std::vector< double >> observed\_counts ()
- · void print observed counts ()
- void print () const

Prints information about the GEESE.

8.28 Geese Class Reference 161

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

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

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

# Information about the model

#### **Parameters**

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

• size\_t nfuns () const noexcept

Number of functions analyzed.

size\_t nnodes () const noexcept

Number of nodes (interior + leaf)

size\_t nleafs () const noexcept

Number of leaf.

size\_t nterms () const

Number of terms included.

size\_t support\_size () const noexcept

Number of unique sets of sufficient stats.

std::vector< size t > nannotations () const noexcept

Number of annotations.

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

Names of the terms in the model.

size\_t parse\_polytomies (bool verb=true, std::vector< size\_t > \*dist=nullptr) const noexcept

Check polytomies and return the largest.

#### Geese prediction

Calculate the conditional probability

#### **Parameters**

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

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

#### Returns

std::vector< double > Returns the posterior probability

- std::vector< std::vector< double >> predict (const std::vector< double > &par, std::vector< std::vector< double >> \*res\_prob=nullptr, bool leave\_one\_out=false, bool only\_annotated=false, bool use\_reduced -\_sequence=true)
- std::vector< std::vector< double > > predict\_backend (const std::vector< double > &par, bool use\_
   reduced\_sequence, const std::vector< size\_t > &preorder)
- std::vector< std::vector< double > > predict\_exhaust\_backend (const std::vector< double > &par, const std::vector< size\_t > &preorder)
- std::vector< std::vector< double > > predict exhaust (const std::vector< double > &par)
- std::vector< std::vector< double > > predict\_sim (const std::vector< double > &par, bool only\_ 
  annotated=false, size t nsims=10000u)

# Non-const pointers to shared objects in <tt>Geese</tt>

These functions provide direct access to some member objects that are shared by the nodes within Geese.

#### Returns

```
get_rengine() returns the Pseudo-RNG engine used.
get_counters() returns the vector of counters used.
get_model() returns the Model object used.
get_support_fun() returns the computed support of the model.
```

- std::mt19937 \* get\_rengine ()
- PhyloCounters \* get\_counters ()
- PhyloModel \* get\_model ()
- PhyloSupport \* get\_support\_fun ()

# **Public Attributes**

- size t nfunctions
- std::map< size\_t, Node > nodes
- barry::MapVec\_type< size\_t > map\_to\_state\_id
- std::vector< std::vector< size\_t >> > pset\_loc

163

Locations of columns.

- std::vector< size\_t > sequence
- std::vector< size t > reduced sequence
- bool initialized = false
- bool delete rengine = false
- bool delete\_support = false

### **Static Public Attributes**

```
    static const size_t etype_default = 1ul
```

- static const size\_t etype\_speciation = 0ul
- static const size\_t etype\_duplication = 1ul
- static const size\_t etype\_either = 2ul

# 8.28.1 Detailed Description

Annotated Phylo Model.

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

Class representing a phylogenetic tree model with annotations.

The Geese class represents a phylogenetic tree model with annotations. It includes a total of N+1 nodes, the +1 being the root node. The class provides methods for initializing the model, calculating the likelihood, simulating trees, and making predictions.

The class includes shared objects within a Geese object, such as rengine, model, states, n\_zeros, n\_ $\leftarrow$  ones, n\_dupl\_events, and n\_spec\_events. It also includes information about the type of event, such as etype\_default, etype\_speciation, etype\_duplication, and etype\_either.

The class provides constructors, a destructor, and methods for initializing the model, inheriting support, calculating the sequence, calculating the 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 103 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

165

### 8.28.3.1 calc\_reduced\_sequence()

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

Definition at line 358 of file geese-meat.hpp.

### 8.28.3.2 calc\_sequence()

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

Definition at line 314 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 480 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 769 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 792 of file geese-meat.hpp.

### 8.28.3.6 get\_counters()

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

Definition at line 752 of file geese-meat.hpp.

### 8.28.3.7 get\_model()

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

Definition at line 757 of file geese-meat.hpp.

# 8.28.3.8 get\_probabilities()

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

Definition at line 406 of file geese-meat.hpp.

# 8.28.3.9 get\_rengine()

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

Definition at line 747 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 765 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 761 of file geese-meat.hpp.

### 8.28.3.12 inherit\_support()

Definition at line 257 of file geese-meat.hpp.

# 8.28.3.13 init()

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

### 8.28.3.14 init\_node()

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

### 8.28.3.15 likelihood()

Definition at line 6 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 471 of file geese-meat.hpp.

# 8.28.3.18 nfuns()

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

Number of functions analyzed.

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

### 8.28.3.19 nleafs()

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

Number of leaf.

Definition at line 441 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 434 of file geese-meat.hpp.

### 8.28.3.21 nterms()

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

Number of terms included.

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

# 8.28.3.22 observed\_counts()

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

Definition at line 522 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 487 of file geese-meat.hpp.

### 8.28.3.26 predict()

Definition at line 287 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 656 of file geese-meat.hpp.

### 8.28.3.32 print\_nodes()

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

Definition at line 674 of file geese-meat.hpp.

### 8.28.3.33 print\_observed\_counts()

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

Definition at line 593 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 461 of file geese-meat.hpp.

### 8.28.3.37 update\_annotations()

Definition at line 285 of file geese-meat.hpp.

### 8.28.4 Member Data Documentation

### 8.28.4.1 delete\_rengine

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

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

# 8.28.4.2 delete\_support

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

Definition at line 143 of file geese-bones.hpp.

# 8.28.4.3 etype\_default

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

Definition at line 156 of file geese-bones.hpp.

### 8.28.4.4 etype\_duplication

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

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

### 8.28.4.5 etype\_either

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

Definition at line 159 of file geese-bones.hpp.

# 8.28.4.6 etype\_speciation

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

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

### 8.28.4.7 initialized

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

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

# 8.28.4.8 map\_to\_state\_id

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

Definition at line 133 of file geese-bones.hpp.

### 8.28.4.9 nfunctions

```
size_t Geese::nfunctions
```

Definition at line 130 of file geese-bones.hpp.

### 8.28.4.10 nodes

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

Definition at line 131 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 134 of file geese-bones.hpp.

### 8.28.4.12 reduced\_sequence

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

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

### 8.28.4.13 sequence

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

Definition at line 137 of file geese-bones.hpp.

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

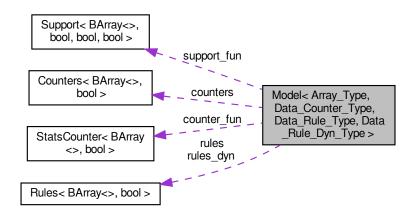
- include/barry/models/geese/geese-bones.hpp
- include/barry/models/geese/geese-meat-constructors.hpp
- include/barry/models/geese/geese-meat-likelihood.hpp
- include/barry/models/geese/geese-meat-likelihood\_exhaust.hpp
- include/barry/models/geese/geese-meat-predict.hpp
- include/barry/models/geese/geese-meat-predict\_exhaust.hpp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

# 8.29 Model < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type > Class Template Reference

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

#include <model-bones.hpp>

Collaboration diagram for Model < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type >:



### **Public Member Functions**

- void set\_rengine (std::mt19937 \*rengine\_, bool delete\_=false)
- void set\_seed (size\_t s)
- Model ()
- Model (size t size )
- Model (const Model< Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type > &Model )
- Model < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type > & operator= (const Model < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type > & Model\_)
- virtual ∼Model ()
- void store\_psets () noexcept
- std::vector< double > gen\_key (const Array\_Type &Array\_)
- size\_t add\_array (const Array\_Type &Array\_, bool force\_new=false)

Adds an array to the support of not already included.

- void print\_stats (size\_t i) const
- · virtual void print () const

Prints information about the model.

- $\bullet \ \, \mathsf{Array\_Type} \ \, \mathsf{8Array\_Type} \ \, \mathsf{\&Array\_Type} \$
- 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 add\_rule\_dyn (Rule\_fun\_type< Array\_Type, Data\_Rule\_Dyn\_Type > count\_fun\_, Data\_Rule\_Dyn
   —Type data\_)
- void set\_rules\_dyn (Rules< Array\_Type, Data\_Rule\_Dyn\_Type > \*rules\_)

#### Likelihood functions.

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

### **Parameters**

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

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

### Extract elements by index

#### **Parameters**

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

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

#### Size of the model

Number of different supports included in the model

This will return the size of stats\_target.

#### Returns

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

- size\_t size () const noexcept
- size\_t size\_unique () const noexcept
- size\_t nterms () const noexcept
- size\_t nrules () const noexcept
- size\_t nrules\_dyn () const noexcept
- size\_t support\_size () const noexcept
- std::vector< std::string > colnames () const
- std::vector< std::vector< double >>\* get\_stats\_target ()

Raw pointers to the support and target statistics.

- std::vector< std::vector< double >> \* get\_stats\_support ()
- std::vector< size\_t > \* get\_arrays2support ()
- std::vector< std::vector< Array\_Type > > \* get\_pset\_arrays ()
- std::vector< std::vector< double >> \* get\_pset\_stats ()

Statistics of the support(s)

- std::vector< std::vector< double >> \* get\_pset\_probs ()
- void set\_transform\_model (std::function< std::vector< double >(double \*, size\_t)> fun, std::vector< std

   ::string > names)

Set the transform\_model\_fun object.

• std::vector< double > transform\_model (double \*data, size\_t k)

### **Protected Attributes**

• MapVec\_type< double, size\_t > keys2support

Map of types of arrays to support sets.

std::vector< std::vector< double >> params\_last

Vector of the previously used parameters.

- std::vector< double > normalizing constants
- std::vector< bool > first\_calc\_done
- bool delete\_counters = false
- bool delete rules = false
- bool delete\_rules\_dyn = false
- std::function< std::vector< double >double \*, size\_t k)> transform\_model\_fun = nullptr

Transformation of the model.

• std::vector< std::string > transform\_model\_term\_names

### Random number generation

Random number generation

- std::mt19937 \* rengine = nullptr
- bool delete\_rengine = false

### Information about the arrays used in the model

stats\_target holds the observed sufficient statistics for each array in the dataset. array\_← frequency contains the frequency with which each of the target stats\_target (arrays) shows in the support. array2support maps array indices (0, 1, ...) to the corresponding support.

Each vector of stats\_support has the data stored in a row-wise order, with each row starting with the weights, e.g., in a model with k terms the first k + 1 elements of stats\_support would be:

- · weights
- term 1
- term 2
- ...
- term k
- std::vector < std::vector < double > > stats\_support
   Sufficient statistics of the model (support)
- std::vector< size t > stats support n arrays

Number of arrays included per support.

 $\bullet \ \, {\sf std::vector} < {\sf std::vector} < {\sf double} > > {\sf stats\_target}$ 

Target statistics of the model.

std::vector< size\_t > arrays2support

# Container space for the powerset (and its sufficient stats\_target)

This is useful in the case of using simulations or evaluating functions that need to account for the full set of states.

- bool with pset = false
- std::vector< std::vector< Array\_Type >> pset\_arrays

Arrays of the support(s)

std::vector< std::vector< double >> pset\_stats

Statistics of the support(s)

std::vector< std::vector< double >> pset\_probs

Probabilities of the support(s)

### Functions to compute statistics

Arguments are recycled to save memory and computation.

- 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 140 of file model-meat.hpp.

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

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

# 8.29.2.3 Model() [3/3]

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

# 8.29.2.4 ∼Model()

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

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

# 8.29.3 Member Function Documentation

# 8.29.3.1 add\_array()

Adds an array to the support of not already included.

# Parameters

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

#### Returns

The number of the array.

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

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

# 8.29.3.4 add\_hasher()

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

#### 8.29.3.6 add rule() [2/2]

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

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

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

## 8.29.3.9 colnames()

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

# 8.29.3.10 conditional\_prob()

Conditional probability ("Gibbs sampler")

Computes the conditional probability of observing  $P\{Y(i,j) = | Y^C, \text{ theta}\}$ , i.e., the probability of observing the entry Y(i,j) equal to one given the rest of the array.

#### **Parameters**

Array←	Array to check
_	
params	Vector of parameters
i	Row entry
j	Column entry

#### Returns

double The conditional probability

# 8.29.3.11 gen\_key()

# 8.29.3.12 get\_arrays2support()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< size_t >* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_\times
Type >::get_arrays2support ()
```

# 8.29.3.13 get\_counters()

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

## 8.29.3.14 get norm const()

# 8.29.3.15 get\_pset()

#### 8.29.3.16 get\_pset\_arrays()

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

## 8.29.3.17 get\_pset\_probs()

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

# 8.29.3.18 get\_pset\_stats() [1/2]

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

Statistics of the support(s)

## 8.29.3.19 get\_pset\_stats() [2/2]

# 8.29.3.20 get\_rengine()

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

## 8.29.3.21 get\_rules()

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

## 8.29.3.22 get\_rules\_dyn()

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

# 8.29.3.23 get\_stats\_support()

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

## 8.29.3.24 get\_stats\_target()

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

Raw pointers to the support and target statistics.

The support of the model is stored as a vector of vector < double>. Each element of it contains the support for an specific type of array included. It represents an array of size  $(k + 1) \times n$  unique elements, with the data stored by-row. The last element of each entry corresponds to the weights, i.e., the frequency with which such sufficient statistics are observed in the support.

# 8.29.3.25 get\_support\_fun()

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

## 8.29.3.26 likelihood() [1/4]

## 8.29.3.27 likelihood() [2/4]

# 8.29.3.28 likelihood() [3/4]

## 8.29.3.29 likelihood() [4/4]

## 8.29.3.30 likelihood total()

# 8.29.3.31 nrules()

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

# 8.29.3.32 nrules\_dyn()

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

# 8.29.3.33 nterms()

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

## 8.29.3.34 operator=()

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

#### 8.29.3.35 print()

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

Prints information about the model.

Definition at line 980 of file model-meat.hpp.

## 8.29.3.36 print\_stats()

# 8.29.3.37 sample() [1/2]

# 8.29.3.38 sample() [2/2]

Definition at line 1107 of file model-meat.hpp.

## 8.29.3.39 set counters()

#### 8.29.3.40 set\_rengine()

Definition at line 129 of file model-bones.hpp.

# 8.29.3.41 set\_rules()

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

## 8.29.3.42 set rules dyn()

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

# 8.29.3.43 set\_seed()

Definition at line 139 of file model-bones.hpp.

# 8.29.3.44 set\_transform\_model()

Set the transform model fun object.

The transform\_model function is used to transform the data

## **Parameters**

data	
target	
n_arrays	
arrays2support	

# 8.29.3.45 size()

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

# 8.29.3.46 size\_unique()

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

## 8.29.3.47 store psets()

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

## 8.29.3.48 support\_size()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::support_←
size ( ) const [noexcept]
```

## 8.29.3.49 transform\_model()

# 8.29.4 Member Data Documentation

# 8.29.4.1 arrays2support

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< size_t > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::arrays2support [protected]
```

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

#### 8.29.4.2 counter\_fun

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\(\cong \)

Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>

StatsCounter<Array_Type, Data_Counter_Type> Model< Array_Type, Data_Counter_Type, Data_Rule_Dyn_Type >::counter_fun [protected]
```

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

#### 8.29.4.3 counters

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Counters<Array_Type, Data_Counter_Type>* Model< Array_Type, Data_Counter_Type, Data_Rule_Dyn_Type >::counters [protected]
```

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

## 8.29.4.4 delete\_counters

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_←
counters = false [protected]
```

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

#### 8.29.4.5 delete rengine

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_←
rengine = false [protected]
```

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

## 8.29.4.6 delete\_rules

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_rules
= false [protected]
```

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

# 8.29.4.7 delete\_rules\_dyn

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_←
rules_dyn = false [protected]
```

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

# 8.29.4.8 first\_calc\_done

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< bool > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type
>::first_calc_done [protected]
```

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

# 8.29.4.9 keys2support

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
MapVec_type< double, size_t > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_←
Rule_Dyn_Type >::keys2support [protected]
```

Map of types of arrays to support sets.

This is of the same length as the vector stats\_target.

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

# 8.29.4.10 normalizing constants

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< double > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::normalizing_constants [protected]
```

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

# 8.29.4.11 params\_last

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector<double> > Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::params_last [protected]
```

Vector of the previously used parameters.

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

## 8.29.4.12 pset\_arrays

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector< Array_Type >> Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::pset_arrays [protected]
```

Arrays of the support(s)

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

## 8.29.4.13 pset probs

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector<double> > Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::pset_probs [protected]
```

Probabilities of the support(s)

Definition at line 83 of file model-bones.hpp.

# 8.29.4.14 pset\_stats

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector<double> > Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::pset_stats [protected]
```

Statistics of the support(s)

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

## 8.29.4.15 rengine

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::mt19937* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >←
::rengine = nullptr [protected]
```

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

#### 8.29.4.16 rules

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Rules<Array_Type, Data_Rule_Type>* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data←
_Rule_Dyn_Type >::rules [protected]
```

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

# 8.29.4.17 rules\_dyn

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Rules<Array_Type, Data_Rule_Dyn_Type>* Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::rules_dyn [protected]
```

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

## 8.29.4.18 stats support

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector< double > > Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::stats_support [protected]
```

Sufficient statistics of the model (support)

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

## 8.29.4.19 stats\_support\_n\_arrays

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< size_t > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::stats_support_n_arrays [protected]
```

Number of arrays included per support.

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

## 8.29.4.20 stats\_target

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector< double > > Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::stats_target [protected]
```

Target statistics of the model.

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

## 8.29.4.21 support fun

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Support<Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type> Model< Array_Type,
Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::support_fun [protected]
```

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

## 8.29.4.22 transform\_model\_fun

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::function<std::vector<double>double *, size_t k)> Model< Array_Type, Data_Counter_Type,
Data_Rule_Type, Data_Rule_Dyn_Type >::transform_model_fun = nullptr [protected]
```

Transformation of the model.

When specified, this function will update the model by modifying the linear equation. For example, if the user wanted to add interaction terms, rescale, or apply other operations of the sorts, the user can do such through this function.

The function should return void and receive the following arguments:

- data Pointer to the first element of the set of sufficient statistics
- k size\_t indicating the number of sufficient statistics

## Returns

Nothing, but it will modify the model data.

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

## 8.29.4.23 transform\_model\_term\_names

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::string > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_\times
Dyn_Type >::transform_model_term_names [protected]
```

Definition at line 125 of file model-bones.hpp.

## 8.29.4.24 with\_pset

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarray_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::with_pset = false [protected]
```

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

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

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

# 8.30 NetCounterData Class Reference

Data class used to store arbitrary size\_t or double vectors.

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

# **Public Member Functions**

- NetCounterData ()
- NetCounterData (const std::vector < size\_t > indices\_, const std::vector < double > numbers\_)
- ∼NetCounterData ()

#### **Public Attributes**

- std::vector< size t > indices
- std::vector< double > numbers

# 8.30.1 Detailed Description

Data class used to store arbitrary size\_t or double vectors.

Definition at line 56 of file network.hpp.

# 8.30.2 Constructor & Destructor Documentation

# 8.30.2.1 NetCounterData() [1/2]

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

Definition at line 62 of file network.hpp.

# 8.30.2.2 NetCounterData() [2/2]

Definition at line 63 of file network.hpp.

# 8.30.2.3 ~NetCounterData()

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

Definition at line 68 of file network.hpp.

## 8.30.3 Member Data Documentation

# 8.30.3.1 indices

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

Definition at line 59 of file network.hpp.

# 8.30.3.2 numbers

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

Definition at line 60 of file network.hpp.

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

• include/barry/counters/network.hpp

# 8.31 NetworkData Class Reference

Data class for Networks.

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

# **Public Member Functions**

- · NetworkData ()
- $\bullet \ \ {\sf NetworkData} \ ({\sf std}::{\sf vector}{<} \ {\sf double} > {\sf vertex\_attr\_}, \ {\sf bool} \ {\sf directed\_=true})$

Constructor using a single attribute.

NetworkData (std::vector< std::vector< double >> vertex\_attr\_, bool directed\_=true)

Constructor using multiple attributes.

∼NetworkData ()

# **Public Attributes**

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

# 8.31.1 Detailed Description

Data class for Networks.

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

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

Definition at line 19 of file network.hpp.

# 8.31.2 Constructor & Destructor Documentation

# 8.31.2.1 NetworkData() [1/3]

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

Definition at line 25 of file network.hpp.

# 8.31.2.2 NetworkData() [2/3]

Constructor using a single attribute.

#### **Parameters**

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

Definition at line 33 of file network.hpp.

# 8.31.2.3 NetworkData() [3/3]

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

Constructor using multiple attributes.

## **Parameters**

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

Definition at line 45 of file network.hpp.

# 8.31.2.4 ~NetworkData()

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

Definition at line 51 of file network.hpp.

# 8.31.3 Member Data Documentation

# 8.31.3.1 directed

bool NetworkData::directed = true

Definition at line 22 of file network.hpp.

8.32 Node Class Reference 201

# 8.31.3.2 vertex\_attr

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

Definition at line 23 of file network.hpp.

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

• include/barry/counters/network.hpp

# 8.32 Node Class Reference

A single node for the model.

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

Collaboration diagram for Node:



# **Public Member Functions**

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

# Construct a new Node object

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

# **Public Attributes**

```
size_t id
```

Id of the node (as specified in the input)

· size tord

Order in which the node was created.

PhyloArray array

Array of the node.

std::vector< size\_t > annotations

Observed annotations (only defined for Geese)

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

Arrays given all possible states.

Node \* parent = nullptr

Parent node.

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

Offspring nodes.

std::vector< size\_t > narray = {}

ID of the array in the model.

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

Induced subtree probabilities.

std::vector< double > probability

The probability of observing each state.

# 8.32.1 Detailed Description

A single node for the model.

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

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

# 8.32.2 Constructor & Destructor Documentation

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

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

Definition at line 37 of file geese-node-bones.hpp.

8.32 Node Class Reference 203

# 8.32.2.2 Node() [2/5]

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

## 8.32.2.3 Node() [3/5]

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

# 8.32.2.4 Node() [4/5]

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

# 8.32.2.5 Node() [5/5]

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

# 8.32.2.6 ∼Node()

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

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

# 8.32.3 Member Function Documentation

# 8.32.3.1 get\_parent()

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

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

# 8.32.3.2 is\_leaf()

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

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

# 8.32.3.3 noffspring()

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

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

## 8.32.4 Member Data Documentation

# 8.32.4.1 annotations

```
std::vector< size_t > Node::annotations
```

Observed annotations (only defined for Geese)

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

## 8.32.4.2 array

PhyloArray Node::array

Array of the node.

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

8.32 Node Class Reference 205

# 8.32.4.3 arrays

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

Arrays given all possible states.

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

# 8.32.4.4 duplication

```
bool Node::duplication
```

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

# 8.32.4.5 id

```
size_t Node::id
```

Id of the node (as specified in the input)

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

# 8.32.4.6 narray

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

ID of the array in the model.

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

# 8.32.4.7 offspring

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

Offspring nodes.

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

# 8.32.4.8 ord

```
size_t Node::ord
```

Order in which the node was created.

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

## 8.32.4.9 parent

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

Parent node.

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

# 8.32.4.10 probability

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

The probability of observing each state.

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

# 8.32.4.11 subtree\_prob

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

Induced subtree probabilities.

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

#### 8.32.4.12 visited

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

Definition at line 26 of file geese-node-bones.hpp.

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

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

# 8.33 NodeData Class Reference

Data definition for the PhyloArray class.

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

# **Public Member Functions**

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

# **Public Attributes**

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

# 8.33.1 Detailed Description

Data definition for the PhyloArray class.

This holds basic information about a given node.

Definition at line 12 of file geese-types.hpp.

# 8.33.2 Constructor & Destructor Documentation

# 8.33.2.1 NodeData()

Definition at line 32 of file geese-types.hpp.

## 8.33.3 Member Data Documentation

# 8.33.3.1 blengths

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

Branch length.

Definition at line 18 of file geese-types.hpp.

# 8.33.3.2 duplication

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

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

# 8.33.3.3 states

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

State of the parent node.

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

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

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

# 8.34 PhyloCounterData Class Reference

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

# **Public Member Functions**

```
• PhyloCounterData (std::vector< size_t > data_, std::vector< double > *counters_=nullptr)
```

- PhyloCounterData ()
- size\_t at (size\_t d)
- size\_t operator() (size\_t d)
- size\_t operator[] (size\_t d)
- void reserve (size\_t x)
- void push\_back (size\_t x)
- void shrink\_to\_fit ()
- size\_t size ()
- std::vector< size\_t >::iterator begin ()
- std::vector< size\_t >::iterator end ()
- bool empty ()
- std::vector< double > \* get\_counters ()

# 8.34.1 Detailed Description

Definition at line 42 of file geese-types.hpp.

# 8.34.2 Constructor & Destructor Documentation

# 8.34.2.1 PhyloCounterData() [1/2]

Definition at line 48 of file geese-types.hpp.

# 8.34.2.2 PhyloCounterData() [2/2]

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

Definition at line 53 of file geese-types.hpp.

# 8.34.3 Member Function Documentation

## 8.34.3.1 at()

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

# 8.34.3.2 begin()

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

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

# 8.34.3.3 empty()

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

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

# 8.34.3.4 end()

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

Definition at line 64 of file geese-types.hpp.

# 8.34.3.5 get\_counters()

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

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

## 8.34.3.6 operator()()

Definition at line 56 of file geese-types.hpp.

# 8.34.3.7 operator[]()

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

# 8.34.3.8 push\_back()

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

# 8.34.3.9 reserve()

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

## 8.34.3.10 shrink\_to\_fit()

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

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

# 8.34.3.11 size()

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

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

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

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

# 8.35 PhyloRuleDynData Class Reference

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

# **Public Member Functions**

- PhyloRuleDynData (const std::vector< double > \*counts\_, size\_t pos\_, size\_t lb\_, size\_t ub\_, size\_
   t duplication\_)
- const double operator() () const
- ∼PhyloRuleDynData ()

# **Public Attributes**

- const std::vector< double > \* counts
- size\_t pos
- size\_t lb
- size\_t ub
- · size t duplication

# 8.35.1 Detailed Description

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

## 8.35.2 Constructor & Destructor Documentation

# 8.35.2.1 PhyloRuleDynData()

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

# 8.35.2.2 ~PhyloRuleDynData()

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

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

# 8.35.3 Member Function Documentation

# 8.35.3.1 operator()()

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

Definition at line 88 of file geese-types.hpp.

# 8.35.4 Member Data Documentation

# 8.35.4.1 counts

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

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

# 8.35.4.2 duplication

size\_t PhyloRuleDynData::duplication

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

## 8.35.4.3 lb

size\_t PhyloRuleDynData::lb

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

# 8.35.4.4 pos

size\_t PhyloRuleDynData::pos

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

## 8.35.4.5 ub

size\_t PhyloRuleDynData::ub

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

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

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

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

Powerset of a binary array.

#include <powerset-bones.hpp>

Collaboration diagram for PowerSet < Array\_Type, Data\_Rule\_Type >:



#### **Public Member Functions**

- void init support ()
- void calc ()
- void reset (size\_t N\_, size\_t M\_)

## Construct and destroy a PowerSet object

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

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

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

- void add\_rule (Rule < Array\_Type, Data\_Rule\_Type > rule)
- void add\_rule (Rule\_fun\_type < Array\_Type, Data\_Rule\_Type > count\_fun\_, Data\_Rule\_Type data\_)

# **Getter functions**

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

# **Public Attributes**

- Array\_Type EmptyArray
- std::vector< Array\_Type > data
- Rules< Array\_Type, Data\_Rule\_Type > \* rules
- size\_t N
- size\_t M
- bool rules deleted = false
- std::vector < size\_t > coordinates\_free
- std::vector< size\_t > coordinates\_locked
- size\_t n\_free
- size\_t n\_locked

# 8.36.1 Detailed Description

template<typename Array\_Type = BArray<>>, typename Data\_Rule\_Type = bool> class PowerSet< Array\_Type, Data\_Rule\_Type >

Powerset of a binary array.

**Template Parameters** 

Array_Type	
Data_Rule_Type	

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

## 8.36.2 Constructor & Destructor Documentation

## 8.36.2.1 PowerSet() [1/3]

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

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

## 8.36.2.2 PowerSet() [2/3]

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

## 8.36.2.3 PowerSet() [3/3]

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

#### 8.36.2.4 ∼PowerSet()

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

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

## 8.36.3 Member Function Documentation

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

Definition at line 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()

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

Definition at line 30 of file progress.hpp.

## 8.37.2.2 ∼Progress()

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

Definition at line 23 of file progress.hpp.

## 8.37.3 Member Function Documentation

#### 8.37.3.1 end()

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

Definition at line 52 of file progress.hpp.

## 8.37.3.2 next()

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

Definition at line 41 of file progress.hpp.

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

• include/barry/progress.hpp

## 8.38 Rule< Array Type, Data Type > Class Template Reference

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

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

## **Public Member Functions**

- ∼Rule ()
- Data\_Type & D ()

Read/Write access to the data.

- bool operator() (const Array\_Type &a, size\_t i, size\_t j)
- std::string & get\_name ()
- std::string & get\_description ()
- std::string get\_name () const
- std::string get\_description () const

## Construct a new Rule object

Construct a new Rule object

#### **Parameters**

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

- Rule ()
- Rule (Rule\_fun\_type< Array\_Type, Data\_Type > fun\_, Data\_Type dat\_, std::string name\_="", std::string desc ="")

## 8.38.1 Detailed Description

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

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

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

#### **Template Parameters**

Array_Type	An object of class BArray.
Data_Type	Any type.

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

## 8.38.2 Constructor & Destructor Documentation

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

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

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

## 8.38.2.2 Rule() [2/2]

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

## 8.38.2.3 ∼Rule()

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

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

#### 8.38.3 Member Function Documentation

## 8.38.3.1 D()

```
template<typename Array_Type , typename Data_Type >
Data_Type & Rule< Array_Type, Data_Type >::D [inline]
```

Read/Write access to the data.

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

## 8.38.3.2 get\_description() [1/2]

```
template<typename Array_Type , typename Data_Type >
std::string & Rule< Array_Type, Data_Type >::get_description [inline]
```

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

## 8.38.3.3 get\_description() [2/2]

```
template<typename Array_Type , typename Data_Type >
std::string Rule< Array_Type, Data_Type >::get_description [inline]
```

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

#### 8.38.3.4 get\_name() [1/2]

```
template<typename Array_Type , typename Data_Type >
std::string & Rule< Array_Type, Data_Type >::get_name [inline]
```

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

#### 8.38.3.5 get\_name() [2/2]

```
template<typename Array_Type , typename Data_Type >
std::string Rule< Array_Type, Data_Type >::get_name [inline]
```

Definition at line 60 of file rules-meat.hpp.

#### 8.38.3.6 operator()()

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

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

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

## 8.39 Rules < Array\_Type, Data\_Type > Class Template Reference

Vector of objects of class Rule.

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

## **Public Member Functions**

- Rules ()
- Rules (const Rules < Array\_Type, Data\_Type > &rules\_)
- Rules< Array\_Type, Data\_Type > operator= (const Rules< Array\_Type, Data\_Type > &rules\_)
- ∼Rules ()
- size\_t size () const noexcept
- bool operator() (const Array\_Type &a, size\_t i, size\_t j)

Check whether a given cell is free or locked.

- void get\_seq (const Array\_Type &a, std::vector< size\_t > \*free, std::vector< size\_t > \*locked=nullptr)
   Computes the sequence of free and locked cells in an BArray.
- std::vector< std::string > get\_names () const
- std::vector< std::string > get\_descriptions () const
- std::vector< Rule< Array\_Type, Data\_Type > >::iterator begin ()
- std::vector< Rule< Array\_Type, Data\_Type > >::iterator end ()

## Rule adding

#### **Parameters**

rule

- void add\_rule (Rule < Array\_Type, Data\_Type > rule)
- void add\_rule (Rule\_fun\_type< Array\_Type, Data\_Type > rule\_, Data\_Type data\_, std::string name\_="", std::string description\_="")

## 8.39.1 Detailed Description

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

Vector of objects of class Rule.

## **Template Parameters**

Array_Type	An object of class BArray
Data_Type	Any type.

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

#### 8.39.2 Constructor & Destructor Documentation

## 8.39.2.1 Rules() [1/2]

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

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

## 8.39.2.2 Rules() [2/2]

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

## 8.39.2.3 ∼Rules()

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

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

#### 8.39.3 Member Function Documentation

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

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

#### 8.39.3.2 add rule() [2/2]

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

## 8.39.3.3 begin()

```
template<typename Array_Type , typename Data_Type >
std::vector< Rule<Array_Type,Data_Type> >::iterator Rules< Array_Type, Data_Type >::begin (
) [inline]
```

Definition at line 134 of file rules-bones.hpp.

## 8.39.3.4 end()

```
template<typename Array_Type , typename Data_Type >
std::vector< Rule<Array_Type,Data_Type> >::iterator Rules< Array_Type, Data_Type >::end ( )
[inline]
```

Definition at line 137 of file rules-bones.hpp.

## 8.39.3.5 get\_descriptions()

```
template<typename Array_Type , typename Data_Type >
std::vector< std::string > Rules< Array_Type, Data_Type >::get_descriptions [inline]
```

Definition at line 180 of file rules-meat.hpp.

#### 8.39.3.6 get\_names()

```
template<typename Array_Type , typename Data_Type >
std::vector< std::string > Rules< Array_Type, Data_Type >::get_names [inline]
```

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

## 8.39.3.7 get\_seq()

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

## **Parameters**

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

#### Returns

Nothing.

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

## 8.39.3.8 operator()()

Check whether a given cell is free or locked.

#### **Parameters**

а	A BArray object
i	row position
j	col position

#### Returns

true If the cell is locked false If the cell is free

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

## 8.39.3.9 operator=()

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

## 8.39.3.10 size()

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

Definition at line 84 of file rules-bones.hpp.

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

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

## 8.40 StatsCounter< Array\_Type, Data\_Type > Class Template Reference

Count stats for a single Array.

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

#### **Public Member Functions**

```
• StatsCounter (const Array_Type *Array_)
```

Creator of a StatsCounter

StatsCounter (const StatsCounter < Array\_Type, Data\_Type > &counter)

Copy constructor.

• StatsCounter ()

Can be created without setting the array.

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

Changes the reference array for the counting.

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

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

- void count\_current (size\_t i, size\_t j)
- std::vector< double > count\_all ()
- Counters< Array Type, Data Type > \* get counters ()
- std::vector< std::string > get\_names () const
- std::vector< std::string > get\_descriptions () const
- size t size () const

## 8.40.1 Detailed Description

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

Count stats for a single Array.

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

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

#### 8.40.2 Constructor & Destructor Documentation

#### 8.40.2.1 StatsCounter() [1/3]

Creator of a StatsCounter

#### **Parameters**

Array←	A const pointer to a BArray.

Definition at line 37 of file statscounter-bones.hpp.

## 8.40.2.2 StatsCounter() [2/3]

Copy constructor.

**Parameters** 

counter

## 8.40.2.3 StatsCounter() [3/3]

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

Can be created without setting the array.

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

## 8.40.2.4 $\sim$ StatsCounter()

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

## 8.40.3 Member Function Documentation

## 8.40.3.1 add\_counter()

## 8.40.3.2 count\_all()

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

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

#### 8.40.3.3 count\_current()

#### 8.40.3.4 count\_init()

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

## 8.40.3.5 get\_counters()

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

## 8.40.3.6 get\_descriptions()

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

#### 8.40.3.7 get\_names()

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

## 8.40.3.8 reset\_array()

Changes the reference array for the counting.

#### **Parameters**

Array⇔	A pointer to an array of class Array_Type.

#### 8.40.3.9 set counters()

#### 8.40.3.10 size()

```
template<typename Array_Type , typename Data_Type >
size_t StatsCounter< Array_Type, Data_Type >::size ( ) const [inline]
```

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

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

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

# 8.41 Support < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type > Class Template Reference

Compute the support of sufficient statistics.

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

## **Public Member Functions**

Support (const Array\_Type &Array\_)

Constructor passing a reference Array.

• Support (size\_t N\_, size\_t M\_)

Constructor specifying the dimensions of the array (empty).

- Support ()
- ∼Support ()
- void init\_support (std::vector< Array\_Type > \*array\_bank=nullptr, std::vector< double > \*stats\_← bank=nullptr)
- void calc (std::vector< Array\_Type > \*array\_bank=nullptr, std::vector< double > \*stats\_bank=nullptr, size
   \_t max\_num\_elements\_=0u)

Computes the entire support.

- std::vector< double > get\_counts () const
- std::vector< double > \* get current stats ()

List current statistics.

- · void print () const
- const 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 416 of file support-meat.hpp.

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

Definition at line 443 of file support-meat.hpp.

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

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

#### 8.41.3.4 add\_rule\_dyn() [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 >::add_rule_\leftrightarrow dyn ( Rule< Array_Type, Data_Rule_Dyn_Type > * f_ ) [inline]
```

Definition at line 478 of file support-meat.hpp.

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

```
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 >::add_rule_\leftrightarrow dyn ( Rule< Array_Type, Data_Rule_Dyn_Type > f_- ) [inline]
```

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

## 8.41.3.6 calc()

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

Computes the entire support.

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

#### **Parameters**

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

Definition at line 382 of file support-meat.hpp.

#### 8.41.3.7 eval\_rules\_dyn()

Definition at line 513 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_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 >
std::vector< double > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_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 426 of file support-meat.hpp.

## 8.41.3.19 set\_rules()

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

#### 8.41.3.20 set\_rules\_dyn()

Definition at line 498 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← _Type >::change_stats
```

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

## 8.41.4.2 coordiantes\_n\_free

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::coordiantes↔
n free
```

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

#### 8.41.4.3 coordiantes\_n\_locked

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::coordiantes \cdot _n_locked
```

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

#### 8.41.4.4 coordinates free

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< size_t > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn 
_Type >::coordinates_free
```

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

#### 8.41.4.5 coordinates\_locked

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< size_t > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn 
_Type >::coordinates_locked
```

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

## 8.41.4.6 current\_stats

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< double > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn← _Type >::current_stats
```

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

## 8.41.4.7 delete\_counters

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_← counters = true
```

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

#### 8.41.4.8 delete\_rules

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_← rules = true
```

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

#### 8.41.4.9 delete rules dyn

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_← rules_dyn = true
```

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

#### 8.41.4.10 hashes

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< size_t > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn← _Type >::hashes
```

Definition at line 83 of file support-bones.hpp.

## 8.41.4.11 hashes\_initialized

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< bool > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_↔
Type >::hashes_initialized
```

Definition at line 84 of file support-bones.hpp.

#### 8.41.4.12 M

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename
Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::M
```

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

#### 8.41.4.13 max\_num\_elements

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename
Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::max_num
_elements = BARRY_MAX_NUM_ELEMENTS
```

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

#### 8.41.4.14 N

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename
Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::N
```

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

#### 8.41.4.15 n counters

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::n_
counters
```

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

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

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

## 8.42 vecHasher < T > Struct Template Reference

#include <typedefs.hpp>

## **Public Member Functions**

• std::size\_t operator() (std::vector< T > const &dat) const noexcept

## 8.42.1 Detailed Description

```
template < typename T> struct vecHasher < T>
```

Definition at line 105 of file typedefs.hpp.

## 8.42.2 Member Function Documentation

## 8.42.2.1 operator()()

Definition at line 108 of file typedefs.hpp.

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

• include/barry/typedefs.hpp

## **Chapter 9**

## **File Documentation**

## 9.1 include/barry/barray-bones.hpp File Reference

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



## Classes

class BArray < Cell\_Type, Data\_Type >
 Baseline class for binary arrays.

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

## Classes

class ConstBArrayRowIter< Cell\_Type, Data\_Type >

246 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

248 File Documentation

## 9.3.2.1 BARRAY\_TEMPLATE() [1/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator* ) const &
```

Definition at line 88 of file barray-meat-operators.hpp.

## 9.3.2.2 BARRAY\_TEMPLATE() [2/6]

## 9.3.2.3 BARRAY\_TEMPLATE() [3/6]

Definition at line 46 of file barray-meat-operators.hpp.

## 9.3.2.4 BARRAY\_TEMPLATE() [4/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator- ) const
```

## 9.3.2.5 BARRAY\_TEMPLATE() [5/6]

Definition at line 75 of file barray-meat-operators.hpp.

## **9.3.2.6 BARRAY\_TEMPLATE()** [6/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator/ ) const &
```

Definition at line 105 of file barray-meat-operators.hpp.

## 9.3.2.7 BARRAY\_TEMPLATE\_ARGS()

```
template BARRAY_TEMPLATE_ARGS ( ) const \&
```

## 9.3.2.8 BARRAY\_TYPE()

```
template const BARRAY_TYPE ( ) &
```

Definition at line 20 of file barray-meat-operators.hpp.

## 9.3.2.9 for()

```
for ( ) [pure virtual]
```

Definition at line 66 of file barray-meat-operators.hpp.

## 9.3.2.10 operator()()

## 9.3.3 Variable Documentation

250 File Documentation

## 9.3.3.1 rhs

```
Data_Type & rhs
Initial value:
{
    checkdim_(*this, rhs)
```

Definition at line 33 of file barray-meat-operators.hpp.

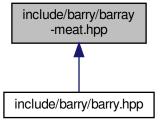
## 9.3.3.2 this

```
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 >

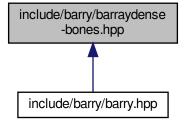
# 9.6 include/barry/barraycell-meat.hpp File Reference

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

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 &
```

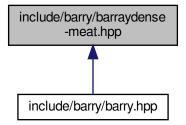
## 9.8.2.6 BDENSE\_TYPE()

```
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( \label{eq:a_b} 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

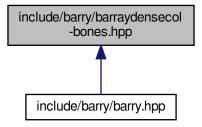
#### 9.11.1.1 POS

```
#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.

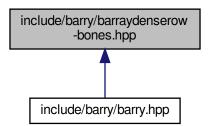
# 9.12.1.3 ZERO\_CELL

```
#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

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

### 9.13.1.2 POS\_N

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

#### 9.13.1.3 ZERO\_CELL

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

Definition at line 6 of file barraydenserow-bones.hpp.

# 9.14 include/barry/barrayrow-bones.hpp File Reference

#### **Classes**

- class BArrayRow
   Cell\_Type, Data\_Type >
- class BArrayRow const< Cell Type, Data Type >

# 9.15 include/barry/barrayrow-meat.hpp File Reference

# **Macros**

- #define BROW\_TYPE() BArrayRow<Cell\_Type, Data\_Type>
- #define BROW\_TEMPLATE\_ARGS() < typename Cell\_Type, typename Data\_Type >
- #define BROW\_TEMPLATE(a, b) template BROW\_TEMPLATE\_ARGS() inline a BROW\_TYPE()::b

### **Functions**

- BROW\_TEMPLATE (void, operator=)(const BROW\_TYPE() &val)
- BROW\_TEMPLATE (void, operator+=)(const BROW\_TYPE() &val)
- BROW\_TEMPLATE (void, operator-=)(const BROW\_TYPE() &val)
- BROW\_TEMPLATE (void, operator\*=)(const BROW\_TYPE() &val)
- BROW\_TEMPLATE (void, operator/=)(const BROW\_TYPE() &val)

# 9.15.1 Macro Definition Documentation

### 9.15.1.1 BROW\_TEMPLATE

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

### 9.15.1.2 BROW\_TEMPLATE\_ARGS

```
#define BROW_TEMPLATE_ARGS() <typename Cell_Type, typename Data_Type>
```

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

## 9.15.1.3 BROW\_TYPE

```
#define BROW_TYPE( ) BArrayRow<Cell_Type, Data_Type>
```

Definition at line 4 of file barrayrow-meat.hpp.

# 9.15.2 Function Documentation

### 9.15.2.1 BROW\_TEMPLATE() [1/5]

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

## 9.15.2.2 BROW\_TEMPLATE() [2/5]

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

## 9.15.2.3 BROW\_TEMPLATE() [3/5]

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

#### 9.15.2.4 BROW\_TEMPLATE() [4/5]

Definition at line 55 of file barrayrow-meat.hpp.

# 9.15.2.5 BROW\_TEMPLATE() [5/5]

Definition at line 11 of file barrayrow-meat.hpp.

# 9.16 include/barry/barrayvector-bones.hpp File Reference

#### **Classes**

```
    class BArrayVector< Cell_Type, Data_Type >
    Row or column of a BArray
```

class BArrayVector\_const< Cell\_Type, Data\_Type >

# 9.17 include/barry/barrayvector-meat.hpp File Reference

# 9.18 include/barry/barry-configuration.hpp File Reference

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



# **Configuration MACROS**

These are mostly related to performance. The definitions follow:

- BARRY\_USE\_UNORDERED\_MAP If specified, then barry is compiled using std::unordered\_map. Otherwise it will use std::map for the arrays.
- BARRY\_USE\_SAFE\_EXP When specified, it will multiply all likelihoods in Model by (1/-100)/(1/-100) so that numerical overflows are avoided.
- BARRY\_USE\_ISFINITE When specified, it will introduce a macro that checks whether the likelihood is finite or not.
- $printf\_barry$  If not specified, will be defined as printf.
- BARRY\_DEBUG\_LEVEL, when defined, will make things verbose.
- #define BARRY\_SAFE\_EXP -100.0
- #define BARRY ISFINITE(a)
- #define BARRY\_CHECK\_SUPPORT(x, maxs)
- #define printf\_barry printf
- #define BARRY\_MAX\_NUM\_ELEMENTS static\_cast< size\_t >(std::numeric\_limits< size\_t >::max() /2u)
- template<typename Ta , typename Tb >
   using Map = std::map< Ta, Tb >

### 9.18.1 Macro Definition Documentation

### 9.18.1.1 BARRY\_CHECK\_SUPPORT

Definition at line 47 of file barry-configuration.hpp.

### 9.18.1.2 BARRY\_ISFINITE

Definition at line 40 of file barry-configuration.hpp.

# 9.18.1.3 BARRY\_MAX\_NUM\_ELEMENTS

```
#define BARRY_MAX_NUM_ELEMENTS static_cast< size_t >(std::numeric_limits< size_t >::max()
/2u)
```

Definition at line 55 of file barry-configuration.hpp.

### 9.18.1.4 BARRY\_SAFE\_EXP

```
#define BARRY_SAFE_EXP -100.0
```

Definition at line 33 of file barry-configuration.hpp.

# 9.18.1.5 printf\_barry

```
#define printf_barry printf
```

Definition at line 51 of file barry-configuration.hpp.

# 9.18.2 Typedef Documentation

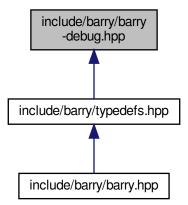
#### 9.18.2.1 Map

```
template<typename Ta , typename Tb >
using Map = std::map<Ta,Tb>
```

Definition at line 27 of file barry-configuration.hpp.

# 9.19 include/barry/barry-debug.hpp File Reference

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



#### **Macros**

• #define BARRY\_DEBUG\_LEVEL 0

## 9.19.1 Macro Definition Documentation

## 9.19.1.1 BARRY\_DEBUG\_LEVEL

#define BARRY\_DEBUG\_LEVEL 0

Definition at line 5 of file barry-debug.hpp.

# 9.20 include/barry/barry-macros.hpp File Reference

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



### **Macros**

- #define BARRY\_ZERO Cell<Cell\_Type>(0.0)
- #define BARRY\_ZERO\_DENSE static\_cast<Cell\_Type>(0.0)
- #define BARRY\_ONE Cell<Cell\_Type>(1.0)
- #define BARRY\_ONE\_DENSE static\_cast<Cell\_Type>(1.0)
- #define BARRY\_UNUSED(expr) do { (void)(expr); } while (0);

#### 9.20.1 Macro Definition Documentation

## 9.20.1.1 BARRY\_ONE

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

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

# 9.20.1.2 BARRY\_ONE\_DENSE

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

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

### 9.20.1.3 BARRY\_UNUSED

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

#### 9.20.1.4 BARRY\_ZERO

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

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

## 9.20.1.5 BARRY\_ZERO\_DENSE

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

Definition at line 5 of file barry-macros.hpp.

# 9.21 include/barry/barry.hpp File Reference

```
#include <iostream>
#include <cstdarg>
#include <vector>
#include <unordered_map>
#include <functional>
#include <stdexcept>
#include <cmath>
#include <map>
#include <algorithm>
#include <utility>
#include <random>
#include <climits>
#include <cfloat>
#include <string>
#include <cstdint>
#include <memory>
#include <regex>
#include <iterator>
#include "typedefs.hpp"
#include "barry-macros.hpp"
#include "freqtable.hpp"
#include "cell-bones.hpp"
#include "cell-meat.hpp"
#include "barray-bones.hpp"
#include "barraycell-bones.hpp"
#include "barray-meat.hpp"
#include "barraycell-meat.hpp"
```

```
#include "barray-meat-operators.hpp"
#include "barraydense-bones.hpp"
#include "barraydensecell-bones.hpp"
#include "barraydenserow-bones.hpp"
#include "barraydensecol-bones.hpp"
#include "barraydense-meat.hpp"
#include "barraydensecell-meat.hpp"
#include "barraydense-meat-operators.hpp"
#include "counters-bones.hpp"
#include "counters-meat.hpp"
#include "statscounter-bones.hpp"
#include "statscounter-meat.hpp"
#include "support-bones.hpp"
#include "support-meat.hpp"
#include "powerset-bones.hpp"
#include "powerset-meat.hpp"
#include "model-bones.hpp"
#include "model-meat.hpp"
#include "rules-bones.hpp"
#include "rules-meat.hpp"
#include "counters/network.hpp"
Include dependency graph for barry.hpp:
```

# **Namespaces**

barry

barry: Your go-to motif accountant

· barry::counters

Tree class and Treelterator class.

· barry::counters::network

#### **Macros**

- #define BARRY\_HPP
- #define BARRY\_VERSION\_MAYOR 0
- #define BARRY\_VERSION\_MINOR 1
- #define BARRY\_VERSION BARRY\_VERSION\_MAYOR ## . ## BARRY\_VERSION\_MINOR
- #define COUNTER FUNCTION(a)
- #define COUNTER\_LAMBDA(a)
- #define RULE\_FUNCTION(a)
- #define RULE LAMBDA(a)

### 9.21.1 Macro Definition Documentation

## 9.21.1.1 BARRY\_HPP

```
#define BARRY_HPP
```

Definition at line 25 of file barry.hpp.

## 9.21.1.2 BARRY\_VERSION

```
#define BARRY_VERSION_BARRY_VERSION_MAYOR ## . ## BARRY_VERSION_MINOR
```

Definition at line 29 of file barry.hpp.

### 9.21.1.3 BARRY\_VERSION\_MAYOR

```
#define BARRY_VERSION_MAYOR 0
```

Definition at line 27 of file barry.hpp.

## 9.21.1.4 BARRY\_VERSION\_MINOR

```
#define BARRY_VERSION_MINOR 1
```

Definition at line 28 of file barry.hpp.

## 9.21.1.5 COUNTER\_FUNCTION

```
#define COUNTER_FUNCTION( a )
```

#### Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
inline double (a) (const Array_Type & Array, size_t i, size_t j, Data_Type & data)\
```

Definition at line 88 of file barry.hpp.

## 9.21.1.6 COUNTER\_LAMBDA

Definition at line 91 of file barry.hpp.

### 9.21.1.7 RULE\_FUNCTION

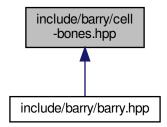
Definition at line 95 of file barry.hpp.

#### 9.21.1.8 RULE LAMBDA

Definition at line 98 of file barry.hpp.

# 9.22 include/barry/cell-bones.hpp File Reference

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

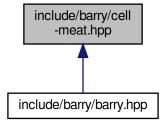


#### **Classes**

class Cell
 Cell\_Type >
 Entries in BArray. For now, it only has two members:

# 9.23 include/barry/cell-meat.hpp File Reference

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



# 9.24 include/barry/col-bones.hpp File Reference

# 9.25 include/barry/counters-bones.hpp File Reference

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

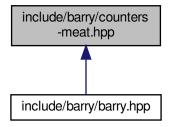


### **Classes**

- class Counter< Array\_Type, Data\_Type >
  - A counter function based on change statistics.
- class Counters
   Array\_Type, Data\_Type >
   Vector of counters.

# 9.26 include/barry/counters-meat.hpp File Reference

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



#### **Macros**

- #define COUNTER\_TYPE() Counter<Array\_Type,Data\_Type>
- #define COUNTER\_TEMPLATE\_ARGS() < typename Array\_Type, typename Data\_Type>
- #define 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\_t i = locator->second
- size\_t size\_t j
- Data\_Type fun
- Data Type counter
- · return
- Data\_Type count\_fun\_
- Data\_Type Counter\_fun\_type< Array\_Type, Data\_Type > init\_fun\_
- Data\_Type Counter\_fun\_type
   Array\_Type, Data\_Type > Hasher\_fun\_type
   Array\_Type, Data\_Type > hasher\_fun\_
- Data\_Type Counter\_fun\_type
   Array\_Type, Data\_Type > Hasher\_fun\_type
   Array\_Type, Data\_Type > Data\_Type data\_
- Data\_Type Counter\_fun\_type
   Array\_Type, Data\_Type > Hasher\_fun\_type
   Array\_Type, Data\_Type > Data\_Type std::string name
- Data\_Type Counter\_fun\_type< Array\_Type, Data\_Type > Hasher\_fun\_type< Array\_Type, Data\_Type > Data\_Type std::string std::string desc\_
- · bool add dims
- · return res
- Data Type fun

### 9.26.1 Macro Definition Documentation

#### 9.26.1.1 COUNTER\_TEMPLATE

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

### 9.26.1.2 COUNTER\_TEMPLATE\_ARGS

```
#define COUNTER_TEMPLATE_ARGS() <typename Array_Type, typename Data_Type>
```

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

### 9.26.1.3 COUNTER\_TYPE

```
#define COUNTER_TYPE( ) Counter<Array_Type,Data_Type>
```

Definition at line 4 of file counters-meat.hpp.

#### 9.26.1.4 COUNTERS\_TEMPLATE

Definition at line 129 of file counters-meat.hpp.

### 9.26.1.5 COUNTERS\_TEMPLATE\_ARGS

```
#define COUNTERS_TEMPLATE_ARGS() <typename Array_Type, typename Data_Type>
```

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

### 9.26.1.6 COUNTERS\_TYPE

```
#define COUNTERS_TYPE( ) Counters<Array_Type,Data_Type>
```

Definition at line 125 of file counters-meat.hpp.

# 9.26.1.7 TMP\_HASHER\_CALL

```
#define TMP_HASHER_CALL Hasher_fun_type<Array_Type,Data_Type>
```

Definition at line 115 of file counters-meat.hpp.

### 9.26.2 Function Documentation

### 9.26.2.1 count fun()

# 9.26.2.2 COUNTER\_TEMPLATE() [1/9]

```
COUNTER_TEMPLATE (

Counter ) const
```

# 9.26.2.3 COUNTER\_TEMPLATE() [2/9]

# 9.26.2.4 COUNTER\_TEMPLATE() [3/9]

# 9.26.2.5 COUNTER\_TEMPLATE() [4/9]

< 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_Rule_Dyn_Type Data_Rule_Dyn_Type data_
Initial value:
{
    rules_dyn->add_rule(
        rule_fun_,
        data_
}
```

Definition at line 182 of file counters-meat.hpp.

### 9.26.3.6 desc\_

```
Data_Type Counter_fun_type<Array_Type,Data_Type> Hasher_fun_type<Array_Type,Data_Type> Data←
_Type std::string std::string desc_
```

#### Initial value:

```
data.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

```
const std::vector< double > size_t i = locator->second
```

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

#### 9.26.3.11 init fun

```
Data_Type Counter_fun_type<Array_Type,Data_Type> init_fun_
```

Definition at line 180 of file counters-meat.hpp.

#### 9.26.3.12 j

```
size_t 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.

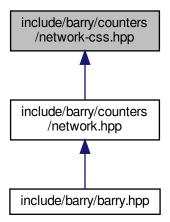
#### 9.26.3.16 return

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.

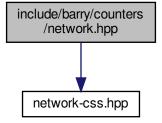
# 9.27.2.15 counter\_css\_partially\_false\_recip\_omiss()

Counts errors of omission.

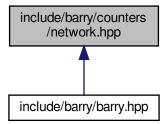
Definition at line 155 of file network-css.hpp.

# 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

#### 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

Function for definition of a network counter function

Definition at line 133 of file network.hpp.

#### 9.28.1.8 NETWORK RULE LAMBDA

Lambda function for definition of a network counter function

Definition at line 138 of file network.hpp.

### 9.28.1.9 NETWORKDENSE\_COUNTER\_LAMBDA

Definition at line 123 of file network.hpp.

# 9.28.2 Typedef Documentation

#### 9.28.2.1 NetCounter

```
template<typename Tnet = Network>
using NetCounter = Counter<Tnet, NetCounterData >
```

Definition at line 89 of file network.hpp.

#### 9.28.2.2 NetCounters

```
template<typename Tnet = Network>
using NetCounters = Counters<Tnet, NetCounterData>
```

Definition at line 92 of file network.hpp.

#### 9.28.2.3 NetModel

```
template<typename Tnet >
using NetModel = Model<Tnet, NetCounterData>
```

Definition at line 101 of file network.hpp.

#### 9.28.2.4 NetRule

```
template<typename Tnet = Network>
using NetRule = Rule<Tnet, bool>
```

Definition at line 104 of file network.hpp.

#### 9.28.2.5 NetRules

```
template<typename Tnet = Network>
using NetRules = Rules<Tnet, bool>
```

Definition at line 107 of file network.hpp.

#### 9.28.2.6 NetStatsCounter

```
template<typename Tnet = Network>
using NetStatsCounter = StatsCounter<Tnet, NetCounterData>
```

Definition at line 98 of file network.hpp.

# 9.28.2.7 NetSupport

```
template<typename Tnet = Network>
using NetSupport = Support<Tnet, NetCounterData >
```

Definition at line 95 of file network.hpp.

#### 9.28.2.8 Network

```
typedef BArray<double, NetworkData> Network
```

Definition at line 82 of file network.hpp.

#### 9.28.2.9 NetworkDense

```
typedef BArrayDense<int, NetworkData> NetworkDense
```

Definition at line 83 of file network.hpp.

# 9.28.3 Function Documentation

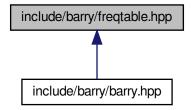
# 9.28.3.1 rules\_zerodiag()

Number of edges.

Definition at line 1381 of file network.hpp.

# 9.29 include/barry/freqtable.hpp File Reference

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



#### **Classes**

class FreqTable
 T >

Frequency table of vectors.

# 9.30 include/barry/model-bones.hpp File Reference

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



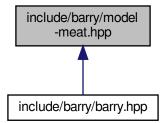
# Classes

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

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

# 9.31 include/barry/model-meat.hpp File Reference

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



#### Macros

- #define MODEL\_TYPE()
- #define MODEL\_TEMPLATE\_ARGS()
- #define MODEL\_TEMPLATE(a, b) template MODEL\_TEMPLATE\_ARGS() inline a MODEL\_TYPE()::b

#### **Functions**

- double update normalizing constant (const double \*params, const double \*support, size t k, size t n)
- double likelihood\_ (const double \*stats\_target, const std::vector< double > &params, const double normalizing constant, size t n params, bool log =false)
- MODEL\_TEMPLATE (void, store\_psets)() noexcept
- MODEL TEMPLATE (std::vector < double >, gen key)(const Array Type &Array )
- MODEL\_TEMPLATE (void, add\_counter)(Counter< Array\_Type</li>
- MODEL\_TEMPLATE (void, set\_counters)(Counters< Array\_Type
- support\_fun set\_counters (counters)
- MODEL TEMPLATE (void, add hasher)(Hasher fun type< Array Type</li>
- MODEL TEMPLATE (void, add rule)(Rule< Array Type</li>
- MODEL\_TEMPLATE (void, set\_rules)(Rules< Array\_Type</li>
- · support fun set rules (rules)
- MODEL\_TEMPLATE (void, add\_rule\_dyn)(Rule < Array\_Type
- MODEL TEMPLATE (void, set rules dyn)(Rules< Array Type</li>
- support fun set rules dyn (rules dyn)
- MODEL\_TEMPLATE (size\_t, add\_array)(const Array\_Type &Array\_
- if (transform\_model\_fun) = transform\_model\_fun(&tmp\_counts[0u], tmp\_counts.size())
- else stats\_target push\_back (counter\_fun.count\_all())
- if (force\_new|(locator==keys2support.end()))
- arrays2support push back (locator->second)
- return arrays2support size () 1u
- MODEL TEMPLATE (double, likelihood)(const std
- MODEL\_TEMPLATE (double, likelihood\_total)(const std
- MODEL\_TEMPLATE (double, get\_norm\_const)(const std
- MODEL\_TEMPLATE (const std::vector< Array\_Type > \*, get\_pset)(const size\_t &i)
- MODEL\_TEMPLATE (const std::vector< double > \*, get\_pset\_stats)(const size\_t &i)
- MODEL\_TEMPLATE (void, print\_stats)(size\_t i) const
- MODEL\_TEMPLATE (size\_t, size)() const noexcept
- MODEL\_TEMPLATE (size\_t, size\_unique)() const noexcept
- MODEL TEMPLATE (size t, nterms)() const noexcept
- MODEL TEMPLATE (size t, nrules)() const noexcept
- MODEL TEMPLATE (size t, nrules dyn)() const noexcept
- MODEL\_TEMPLATE (size\_t, support\_size)() const noexcept
- MODEL\_TEMPLATE (std::vector< std::string >, colnames)() const
- MODEL\_TEMPLATE (Array\_Type, sample)(const Array\_Type &Array\_
- if (locator==keys2support.end())
- std::uniform real distribution urand (0, 1)
- if ((probs.size() > 0u) &&(vec\_equal\_approx(params, params\_last[a])))
- temp\_stats reserve (params.size())
- for (size t array=0u;array< probs.size();++array)</li>
- MODEL\_TEMPLATE (double, conditional\_prob)(const Array\_Type & Array\_
- A insert\_cell (i, j, A.default\_val(), true, false)
- tmp\_counts reserve (counters->size())
- for (size\_t ii=0u;ii < counters->size();++ii) tmp\_counts.push\_back(counters -> operator[](ii).count(A, i, j))
- return (1.0+std::exp(-vec\_inner\_prod< double >(&params[0u], &tmp\_counts[0u], params.size())))
- MODEL\_TEMPLATE (const std::mt19937 \*, get\_rengine)() const

MODEL\_TEMPLATE (std::vector< std::vector< double >> \*, get\_stats\_target)()

```
    MODEL TEMPLATE (std::vector< std::vector< double >> *, get stats support)()

    MODEL_TEMPLATE (std::vector< size_t > *, get_arrays2support)()

    MODEL_TEMPLATE (std::vector< std::vector< Array_Type >> *, get_pset_arrays)()

    MODEL TEMPLATE (std::vector< std::vector< double >> *, get_pset_stats)()

   • MODEL_TEMPLATE (std::vector< std::vector< double >> *, get_pset_probs)()

    MODEL_TEMPLATE (void, set_transform_model)(std

Variables

    Data_Counter_Type & counter

    · return

    Data Counter Type count fun

    Data_Counter_Type Counter_fun_type< Array_Type, Data_Counter_Type > init_fun_

    Data_Counter_Type Counter_fun_type < Array_Type, Data_Counter_Type > Data_Counter_Type data_

    Data_Counter_Type * counters_

    Data_Counter_Type fun_

    Data_Rule_Type & rules

    Data Rule Type * rules

    • this delete rules = false

    Data Rule Dyn Type rule fun

    this rules_dyn = rules_
    • this delete_rules_dyn = false
    · bool force new

    std::vector< double > key = counters->gen_hash(Array_)

    MapVec type< double, size t >::const iterator locator = keys2support.find(key)

    stats_support_n_arrays [locator->second]

    const std::vector< double > & params

    • size_t i = locator->second
    size t a = arrays2support[i]
    double r = urand(*rengine)
    • double cumprob = 0.0
    size_t k = params.size()
    • size t | = 0u
    std::vector< double > & probs = pset_probs[a]

    std::vector< double > temp stats

    const std::vector< double > & stats = pset_stats[a]

    • int i matches = -1
   • return this pset_arrays [a][j]
    • std::vector< double > tmp counts
```

#### 9.31.1 Macro Definition Documentation

template Data\_Counter\_Typetemplate Data\_Rule\_Type

# 9.31.1.1 MODEL\_TEMPLATE

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

#### 9.31.1.2 MODEL TEMPLATE ARGS

Definition at line 128 of file model-meat.hpp.

#### 9.31.1.3 MODEL\_TYPE

Data\_Rule\_Dyn\_Type>

```
template Data_Rule_Dyn_Type * MODEL_TYPE( )

Value:
    Model<Array_Type, Data_Counter_Type, Data_Rule_Type, \)</pre>
```

Definition at line 125 of file model-meat.hpp.

# 9.31.2 Function Documentation

# 9.31.2.1 for() [1/2]

```
for ( )
```

Definition at line 1317 of file model-meat.hpp.

# 9.31.2.2 for() [2/2]

# 9.31.2.3 if() [1/4]

```
if (
     (probs.size() > Ou) &&(vec_equal_approx(params, params_last[a])) )
```

Definition at line 1299 of file model-meat.hpp.

# 9.31.2.4 if() [2/4]

```
if (
     force_new| locator==keys2support.end() )
```

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

Definition at line 457 of file model-meat.hpp.

#### 9.31.2.5 if() [3/4]

```
if (
    locator = = keys2support.end() )
```

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

Definition at line 1201 of file model-meat.hpp.

# 9.31.2.6 if() [4/4]

```
if (
    transform_model_fun ) = transform_model_fun(&tmp_counts[0u], tmp_counts.size())
```

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

# 9.31.2.7 insert\_cell()

# 9.31.2.8 likelihood\_()

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

# 9.31.2.9 MODEL\_TEMPLATE() [1/33]

#### 9.31.2.10 MODEL\_TEMPLATE() [2/33]

Definition at line 1387 of file model-meat.hpp.

# 9.31.2.11 MODEL\_TEMPLATE() [3/33]

Definition at line 927 of file model-meat.hpp.

### 9.31.2.12 MODEL\_TEMPLATE() [4/33]

Definition at line 939 of file model-meat.hpp.

# **9.31.2.13** MODEL\_TEMPLATE() [5/33]

# **9.31.2.14** MODEL\_TEMPLATE() [6/33]

Definition at line 891 of file model-meat.hpp.

# **9.31.2.15 MODEL\_TEMPLATE()** [7/33]

Definition at line 569 of file model-meat.hpp.

# **9.31.2.16** MODEL\_TEMPLATE() [8/33]

Definition at line 825 of file model-meat.hpp.

#### 9.31.2.17 MODEL\_TEMPLATE() [9/33]

# 9.31.2.18 MODEL\_TEMPLATE() [10/33]

Definition at line 1065 of file model-meat.hpp.

# 9.31.2.19 MODEL\_TEMPLATE() [11/33]

Definition at line 1072 of file model-meat.hpp.

#### 9.31.2.20 MODEL\_TEMPLATE() [12/33]

Definition at line 1055 of file model-meat.hpp.

### 9.31.2.21 MODEL\_TEMPLATE() [13/33]

Definition at line 1040 of file model-meat.hpp.

# **9.31.2.22** MODEL\_TEMPLATE() [14/33]

Definition at line 1047 of file model-meat.hpp.

#### 9.31.2.23 MODEL\_TEMPLATE() [15/33]

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

# 9.31.2.24 MODEL\_TEMPLATE() [16/33]

```
MODEL_TEMPLATE (
          std::vector< double > ,
           gen_key ) const &
```

Definition at line 312 of file model-meat.hpp.

#### 9.31.2.25 MODEL\_TEMPLATE() [17/33]

Definition at line 1422 of file model-meat.hpp.

# 9.31.2.26 MODEL\_TEMPLATE() [18/33]

```
MODEL_TEMPLATE (
          std::vector< std::string > ,
          colnames ) const
```

Definition at line 1091 of file model-meat.hpp.

# 9.31.2.27 MODEL\_TEMPLATE() [19/33]

```
MODEL_TEMPLATE (
          std::vector< std::vector< Array_Type > > * ,
          get_pset_arrays )
```

Definition at line 1427 of file model-meat.hpp.

# 9.31.2.28 MODEL\_TEMPLATE() [20/33]

```
MODEL_TEMPLATE (  \mbox{std::vector} < \mbox{std::vector} < \mbox{double} >> * \mbox{,} \\ \mbox{get\_pset\_probs} \mbox{ )}
```

Definition at line 1435 of file model-meat.hpp.

# 9.31.2.29 MODEL\_TEMPLATE() [21/33]

```
MODEL_TEMPLATE (
          std::vector< std::vector< double > > * ,
          get_pset_stats )
```

Definition at line 1431 of file model-meat.hpp.

#### 9.31.2.30 MODEL\_TEMPLATE() [22/33]

```
MODEL_TEMPLATE (
          std::vector< std::vector< double > > * ,
          get_stats_support )
```

Definition at line 1417 of file model-meat.hpp.

### 9.31.2.31 MODEL\_TEMPLATE() [23/33]

```
MODEL_TEMPLATE (
          std::vector< std::vector< double > > * ,
          get_stats_target )
```

Definition at line 1412 of file model-meat.hpp.

# 9.31.2.32 MODEL\_TEMPLATE() [24/33]

# 9.31.2.33 MODEL\_TEMPLATE() [25/33]

# 9.31.2.34 MODEL\_TEMPLATE() [26/33]

#### 9.31.2.35 MODEL\_TEMPLATE() [27/33]

#### 9.31.2.36 MODEL\_TEMPLATE() [28/33]

Definition at line 950 of file model-meat.hpp.

# **9.31.2.37** MODEL\_TEMPLATE() [29/33]

# 9.31.2.38 MODEL\_TEMPLATE() [30/33]

# 9.31.2.39 MODEL\_TEMPLATE() [31/33]

# 9.31.2.40 MODEL\_TEMPLATE() [32/33]

Definition at line 1439 of file model-meat.hpp.

### 9.31.2.41 MODEL\_TEMPLATE() [33/33]

Definition at line 305 of file model-meat.hpp.

# 9.31.2.42 push\_back() [1/2]

# 9.31.2.43 push\_back() [2/2]

#### 9.31.2.44 reserve() [1/2]

# 9.31.2.45 reserve() [2/2]

# 9.31.2.46 return()

```
return (  1.0+ \ std::exp-vec\_inner\_prod < \ double > (\&params[0u], \ \&tmp\_counts[0u], \ params. \leftrightarrow size()) \ )
```

### 9.31.2.47 set\_counters()

# 9.31.2.48 set\_rules()

```
support_fun set_rules (
    rules )
```

# 9.31.2.49 set\_rules\_dyn()

### 9.31.2.50 size()

```
return arrays2support size ( )
```

# 9.31.2.51 update\_normalizing\_constant()

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

#### 9.31.2.52 urand()

```
std::uniform_real_distribution urand ( \begin{array}{c} 0 \text{ ,} \\ 1 \end{array} )
```

#### 9.31.3 Variable Documentation

#### 9.31.3.1 a

```
size_t a = arrays2support[i]
```

Definition at line 1287 of file model-meat.hpp.

# 9.31.3.2 count\_fun\_

```
Data_Counter_Type count_fun_
```

Definition at line 327 of file model-meat.hpp.

#### 9.31.3.3 counter

```
Data_Counter_Type& counter

Initial value:
{
    counters->add_counter(counter, Data_Counter_Type())
```

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

#### 9.31.3.4 counters\_

```
Data_Counter_Type* counters_
Initial value:
{
    if (delete_counters) {
        delete counters;
        delete_counters = false;
    }
    this->counters = counters_
```

Definition at line 343 of file model-meat.hpp.

#### 9.31.3.5 cumprob

```
double cumprob = 0.0
```

Definition at line 1292 of file model-meat.hpp.

### 9.31.3.6 data\_

```
Data_Rule_Dyn_Type Data_Rule_Dyn_Type data_
Initial value:
{
    counters->add_counter(
        count_fun_,
        init_fun_,
        data_
```

Definition at line 329 of file model-meat.hpp.

# 9.31.3.7 Data\_Counter\_Type

```
template Data_Counter_Type
```

Definition at line 1407 of file model-meat.hpp.

# 9.31.3.8 Data\_Rule\_Type

```
template Data_Rule_Type
```

Definition at line 1407 of file model-meat.hpp.

# 9.31.3.9 delete\_rules

```
this delete_rules = false
```

Definition at line 386 of file model-meat.hpp.

#### 9.31.3.10 delete\_rules\_dyn

```
this delete_rules_dyn = false
```

Definition at line 425 of file model-meat.hpp.

#### 9.31.3.11 else

```
else
```

#### Initial value:

```
probs.resize(pset_arrays[a].size())
```

Definition at line 1309 of file model-meat.hpp.

# 9.31.3.12 force\_new

```
bool force_new

Initial value:
{
    counter_fun.reset_array(&Array_)
```

Definition at line 436 of file model-meat.hpp.

# 9.31.3.13 fun\_

```
Data_Counter_Type fun_
Initial value:
{
    counters->add_hash(fun_)
```

Definition at line 360 of file model-meat.hpp.

#### 9.31.3.14 i

```
const std::vector< double > size_t i = locator->second
```

Definition at line 1195 of file model-meat.hpp.

#### 9.31.3.15 i\_matches

```
int i_matches = -1
```

Definition at line 1316 of file model-meat.hpp.

#### 9.31.3.16 init\_fun\_

```
Data_Counter_Type Counter_fun_type<Array_Type,Data_Counter_Type> init_fun_
```

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

# 9.31.3.17 j

```
const std::vector< double > size_t size_t j = 0u
```

Definition at line 1297 of file model-meat.hpp.

# 9.31.3.18 k

```
size_t k = params.size()
```

Definition at line 1294 of file model-meat.hpp.

#### 9.31.3.19 key

```
std::vector< double > key = counters->gen_hash(Array_)
```

Definition at line 455 of file model-meat.hpp.

#### 9.31.3.20 locator

```
MapVec_type< double, size_t >::const_iterator locator = keys2support.find(key)
```

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

# 9.31.3.21 params

```
const std::vector< double > & params

Initial value:
{
    if (!this->with_pset)
        throw std::logic_error("Sampling is only available when store_pset() is active.")
```

Definition at line 1188 of file model-meat.hpp.

#### 9.31.3.22 probs

```
std::vector< double >& probs = pset_probs[a]
```

Definition at line 1298 of file model-meat.hpp.

### 9.31.3.23 pset\_arrays

```
return this pset_arrays[a][j]
```

Definition at line 1350 of file model-meat.hpp.

### 9.31.3.24 r

```
double r = urand(*rengine)
```

Definition at line 1291 of file model-meat.hpp.

#### 9.31.3.25 return

return

Definition at line 323 of file model-meat.hpp.

# 9.31.3.26 rule\_fun\_

```
Data_Rule_Dyn_Type rule_fun_
```

Definition at line 404 of file model-meat.hpp.

#### 9.31.3.27 rules

```
Initial value:
{
    rules->add_rule(rules, Data_Rule_Type())
```

Definition at line 370 of file model-meat.hpp.

# 9.31.3.28 rules\_

```
Data_Rule_Dyn_Type * rules_
Initial value:
{
    if (delete_rules)
        delete rules
```

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

#### 9.31.3.29 rules\_dyn

```
this rules_dyn = rules_
```

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

#### 9.31.3.30 stats

```
const std::vector< double >& stats = pset_stats[a]
```

Definition at line 1314 of file model-meat.hpp.

# 9.31.3.31 stats\_support\_n\_arrays

```
stats_support_n_arrays[locator->second]
```

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

#### 9.31.3.32 temp\_stats

```
std::vector< double > temp_stats
```

Definition at line 1312 of file model-meat.hpp.

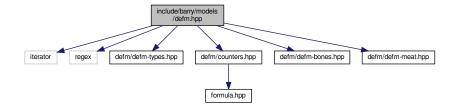
#### 9.31.3.33 tmp\_counts

```
std::vector< double > tmp_counts
```

Definition at line 1369 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:
```

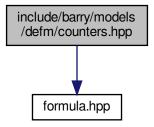


#### **Namespaces**

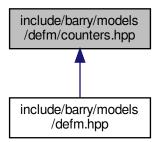
• defm

# 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)
- 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.

#### 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

Lambda function for definition of a network counter function

Definition at line 43 of file counters.hpp.

#### 9.33.1.3 DEFM RULE

Function for definition of a network counter function

Definition at line 53 of file counters.hpp.

#### 9.33.1.4 DEFM\_RULE\_LAMBDA

Lambda function for definition of a network counter function

Definition at line 57 of file counters.hpp.

#### 9.33.1.5 DEFM\_RULEDYN\_LAMBDA

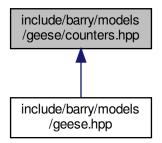
Lambda function for definition of a network counter function

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\_dyn\_limit\_changes (PhyloSupport \*support, size\_t pos, size\_t lb, size\_t ub, size\_
 t duplication=Geese::etype\_default)

Overall functional gains.

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

Extension of a simple counter.

- #define PHYLO\_RULE\_DYN\_LAMBDA(a)
- #define PHYLO\_CHECK\_MISSING()

- std::string get last name (size t d)
- void counter\_overall\_gains (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)
   Overall functional gains.

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

- Functional gains for a specific function (nfun).

   void counter gains k offenring (PhyloCounters \*counters std::vector< size t > nfun size t k=1u size t
- void counter\_gains\_k\_offspring (PhyloCounters \*counters, std::vector< size\_t > nfun, size\_t k=1u, size\_t duplication=Geese::etype default)

k genes gain function nfun

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

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

Keeps track of how many pairs of genes preserve pseudostate.

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

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

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

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

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

  Co-evolution (joint gain or loss)
- void counter\_longest (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

Longest branch mutates (either by gain or by loss)

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

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

  Total number of neofunctionalization events sum\_u sum\_{{w < u}} [x(u,a)\*(1 x(w,a)) + (1 x(u,a)) \* x(w,a)] change

  stat: delta{x(u,a): 0->1} = 1 2 \* x(w,a)
- void counter\_neofun\_a2b (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

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

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

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

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

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

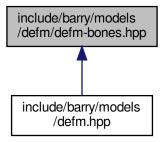
Used when all the functions are in 0 (like the root node prob.)

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

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

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

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

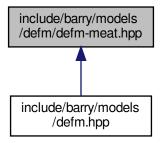


#### **Classes**

· class DEFM

# 9.36 include/barry/models/defm/defm-meat.hpp File Reference

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



#### **Macros**

- #define DEFM\_RANGES(a)
- #define DEFM\_LOOP\_ARRAYS(a) for (size\_t a = 0u; a < (nobs\_i M\_order); ++a)

#### **Functions**

• std::vector< double > keygen\_defm (const DEFMArray &Array\_, DEFMCounterData \*data)

#### 9.36.1 Macro Definition Documentation

#### 9.36.1.1 DEFM\_LOOP\_ARRAYS

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

## 9.36.1.2 DEFM\_RANGES

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

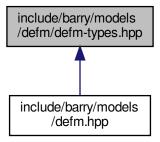
## 9.36.2 Function Documentation

#### 9.36.2.1 keygen\_defm()

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

# 9.37 include/barry/models/defm/defm-types.hpp File Reference

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



#### **Classes**

- class DEFMData
  - Data class for DEFM arrays.
- class DEFMCounterData

Data class used to store arbitrary size t or double vectors.

- class DEFMRuleData
- · class DEFMRuleDynData

# **Typedefs**

typedef barry::BArrayDense< int, DEFMData > DEFMArray

#### Convenient typedefs for network objects.

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

## 9.37.1 Typedef Documentation

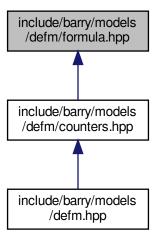
#### 9.37.1.1 **DEFMArray**

typedef barry::BArrayDense<int, DEFMData> DEFMArray

Definition at line 5 of file defm-types.hpp.

# 9.38 include/barry/models/defm/formula.hpp File Reference

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



## **Functions**

void defm\_motif\_parser (std::string formula, std::vector< size\_t > &locations, std::vector< bool > &signs, size t m order, size t y ncol)

Parses a motif formula.

#### 9.38.1 Function Documentation

#### 9.38.1.1 defm\_motif\_parser()

Parses a motif formula.

This function will take the formula and generate the corresponding input for defm::counter\_transition(). Formulas can be specified in the following ways:

- Intercept effect: {...} No transition, only including the current state.
- Transition effect: {...} > {...} Includes current and previous states.

The general notation is  $[0]y[column id]_[row id]$ . A preceeding zero means that the value of the cell is considered to be zero. The column id goes between 0 and the number of columns in the array - 1 (so it is indexed from 0,) and the row id goes from 0 to m\_order.

#### Intercept effects

Intercept effects only involve a single set of curly brackets. Using the 'greater-than' symbol (i.e., '<') is only for transition effects. When specifying intercept effects, users can skip the  $row_id$ , e.g.,  $y0_0$  is equivalent to y0. If the passed  $row_id$  is different from the Markov order, i.e.,  $row_id$  !=  $m_order$ , then the function returns with an error.

#### Examples:

• " $\{y0, 0y1\}$ " is equivalent to set a motif with the first element equal to one and the second to zero.

#### **Transition effects**

Transition effects can be specified using two sets of curly brackets and an greater-than symbol, i.e.,  $\{\ldots\}$  >  $\{\ldots\}$ . The first set of brackets, which we call LHS, can only hold row id that are less than m\_order.

#### **Parameters**

formula	
locations	
signs	
m_order	
y_ncol	

Definition at line 46 of file formula.hpp.

# 9.39 include/barry/models/geese.hpp File Reference

```
#include "geese/geese-types.hpp"
#include "geese/geese-node-bones.hpp"
#include "geese/geese-bones.hpp"
#include "geese/geese-meat.hpp"
#include "geese/geese-meat-constructors.hpp"
#include "geese/geese-meat-likelihood.hpp"
#include "geese/geese-meat-likelihood_exhaust.hpp"
#include "geese/geese-meat-simulate.hpp"
#include "geese/geese-meat-predict.hpp"
#include "geese/geese-meat-predict_exhaust.hpp"
#include "geese/geese-meat-predict_sim.hpp"
#include "geese/flock-bones.hpp"
#include "geese/flock-meat.hpp"
#include "geese/counters.hpp"
#include dependency graph for geese.hpp:
```



## **Namespaces**

• geese

# 9.40 include/barry/models/geese/flock-bones.hpp File Reference

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



## **Classes**

· class Flock

A Flock is a group of Geese.

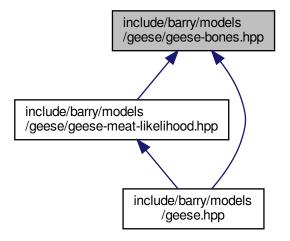
# 9.41 include/barry/models/geese/flock-meat.hpp File Reference

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



# 9.42 include/barry/models/geese/geese-bones.hpp File Reference

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



#### Classes

· class Geese

Annotated Phylo Model.

#### **Macros**

• #define INITIALIZED()

#### **Functions**

```
    template < typename Ta , typename Tb > std::vector < Ta > vector_caster (const std::vector < Tb > &x)
    RULE_FUNCTION (rule_empty_free)
    std::vector < double > keygen_full (const PhyloArray & array, const PhyloCounterData *d)
```

bool vec diff (const std::vector< size t > &s, const std::vector< size t > &a)

## 9.42.1 Macro Definition Documentation

#### **9.42.1.1 INITIALIZED**

```
#define INITIALIZED( )

Value:
    if (!this->initialized) \
    throw std::logic_error("The model has not been initialized yet.");
```

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

#### 9.42.2 Function Documentation

## 9.42.2.1 keygen\_full()

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

#### 9.42.2.2 RULE\_FUNCTION()

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

## 9.42.2.3 vec\_diff()

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

## 9.42.2.4 vector\_caster()

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

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



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

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



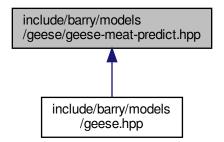


# 9.45 include/barry/models/geese/geese-meat-likelihood\_exhaust.hpp File Reference

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

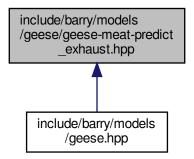


# 9.46 include/barry/models/geese/geese-meat-predict.hpp File Reference



# 9.47 include/barry/models/geese/geese-meat-predict\_exhaust.hpp File Reference

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



# 9.48 include/barry/models/geese/geese-meat-predict\_sim.hpp File Reference

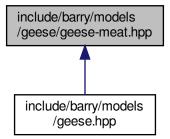


# 9.49 include/barry/models/geese/geese-meat-simulate.hpp File Reference

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

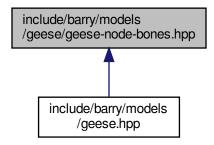


# 9.50 include/barry/models/geese/geese-meat.hpp File Reference



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

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



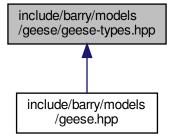
#### **Classes**

· class Node

A single node for the model.

# 9.52 include/barry/models/geese/geese-types.hpp File Reference

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



#### **Classes**

class NodeData

Data definition for the PhyloArray class.

- class PhyloCounterData
- class PhyloRuleDynData

## **Typedefs**

#### Convenient typedefs for Node objects.

- typedef std::vector< std::pair< size\_t, size\_t >> PhyloRuleData
- typedef barry::BArrayDense< size\_t, NodeData > PhyloArray
- typedef barry::Counter< PhyloArray, PhyloCounterData > PhyloCounter
- typedef barry::Counters < PhyloArray, PhyloCounterData > PhyloCounters
- typedef barry::Rule< PhyloArray, PhyloRuleData > PhyloRule
- typedef barry::Rules
   PhyloArray, PhyloRuleData
   PhyloRules
- typedef barry::Rule < PhyloArray, PhyloRuleDynData > PhyloRuleDyn
- typedef barry::Rules
   PhyloArray, PhyloRuleDynData
   PhyloRulesDyn
- typedef barry::Support
   PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloSupport
- typedef barry::StatsCounter< PhyloArray, PhyloCounterData > PhyloStatsCounter
- typedef barry::Model < PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloModel
- typedef barry::PowerSet
   PhyloArray, PhyloRuleData > PhyloPowerSet

## 9.52.1 Typedef Documentation

#### 9.52.1.1 PhyloArray

typedef barry::BArrayDense<size\_t, NodeData> PhyloArray

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

#### 9.52.1.2 PhyloCounter

typedef barry::Counter<PhyloArray, PhyloCounterData > PhyloCounter

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

#### 9.52.1.3 PhyloCounters

typedef barry::Counters< PhyloArray, PhyloCounterData> PhyloCounters

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

#### 9.52.1.4 PhyloModel

typedef barry::Model<PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloModel

Definition at line 113 of file geese-types.hpp.

#### 9.52.1.5 PhyloPowerSet

typedef barry::PowerSet<PhyloArray, PhyloRuleData> PhyloPowerSet

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

#### 9.52.1.6 PhyloRule

typedef barry::Rule<PhyloArray,PhyloRuleData> PhyloRule

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

#### 9.52.1.7 PhyloRuleData

typedef std::vector< std::pair< size\_t, size\_t > > PhyloRuleData

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

## 9.52.1.8 PhyloRuleDyn

typedef barry::Rule<PhyloArray,PhyloRuleDynData> PhyloRuleDyn

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

#### 9.52.1.9 PhyloRules

typedef barry::Rules<PhyloArray,PhyloRuleData> PhyloRules

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

#### 9.52.1.10 PhyloRulesDyn

typedef barry::Rules<PhyloArray,PhyloRuleDynData> PhyloRulesDyn

Definition at line 109 of file geese-types.hpp.

## 9.52.1.11 PhyloStatsCounter

typedef barry::StatsCounter<PhyloArray, PhyloCounterData> PhyloStatsCounter

Definition at line 112 of file geese-types.hpp.

## 9.52.1.12 PhyloSupport

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

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

# 9.53 include/barry/powerset-bones.hpp File Reference

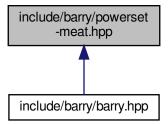
This graph shows which files directly or indirectly include this file:



#### **Classes**

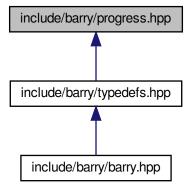
# 9.54 include/barry/powerset-meat.hpp File Reference

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



# 9.55 include/barry/progress.hpp File Reference

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



## **Classes**

• class Progress

A simple progress bar.

## **Macros**

• #define BARRY\_PROGRESS\_BAR\_WIDTH 80

#### 9.55.1 Macro Definition Documentation

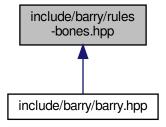
#### 9.55.1.1 BARRY PROGRESS BAR WIDTH

```
#define BARRY_PROGRESS_BAR_WIDTH 80
```

Definition at line 5 of file progress.hpp.

# 9.56 include/barry/rules-bones.hpp File Reference

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



## **Classes**

- class Rule < Array\_Type, Data\_Type >
   Rule for determining if a cell should be included in a sequence.
- class Rules< Array\_Type, Data\_Type >

Vector of objects of class Rule.

## **Functions**

template<typename Array\_Type , typename Data\_Type >
 bool rule\_fun\_default (const Array\_Type \*array, size\_t i, size\_t j, Data\_Type \*dat)

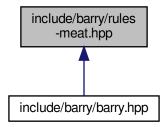
## 9.56.1 Function Documentation

#### 9.56.1.1 rule\_fun\_default()

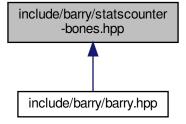
Definition at line 5 of file rules-bones.hpp.

## 9.57 include/barry/rules-meat.hpp File Reference

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



# 9.58 include/barry/statscounter-bones.hpp File Reference



#### Classes

 $\bullet \ \ {\it class StatsCounter} < {\it Array\_Type}, \ {\it Data\_Type} >$ 

Count stats for a single Array.

# 9.59 include/barry/statscounter-meat.hpp File Reference

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



#### **Macros**

- #define STATSCOUNTER TYPE() StatsCounter<Array Type, Data Type>
- #define STATSCOUNTER\_TEMPLATE\_ARGS() < typename Array\_Type, typename Data\_Type >
- #define STATSCOUNTER\_TEMPLATE(a, b) template STATSCOUNTER\_TEMPLATE\_ARGS() inline a STATSCOUNTER\_TYPE()::b

#### **Functions**

- STATSCOUNTER\_TEMPLATE (, StatsCounter)(const StatsCounter< Array\_Type
- EmptyArray clear ()
- STATSCOUNTER TEMPLATE (,~StatsCounter)()
- STATSCOUNTER\_TEMPLATE (void, reset\_array)(const Array\_Type \*Array\_)
- STATSCOUNTER\_TEMPLATE (void, add\_counter)(Counter< Array\_Type</li>
- STATSCOUNTER\_TEMPLATE (void, set\_counters)(Counters< Array\_Type
- STATSCOUNTER\_TEMPLATE (void, count\_init)(size\_t i
- current\_stats resize (counters->size(), 0.0)
- for (size\_t n=0u;n< counters->size();++n) current\_stats[n]
- STATSCOUNTER TEMPLATE (void, count current)(size ti
- STATSCOUNTER\_TEMPLATE (std::vector< std::string >, get\_names)() const
- STATSCOUNTER\_TEMPLATE (std::vector< std::string >, get\_descriptions)() const

#### **Variables**

size\_t j

```
    Data_Type & counter
    EmptyArray = *Array
    current_stats = counter.current_stats
    counters = new Counters<Array_Type,Data_Type>((*counter.counters))
    counter_deleted = false
    Data_Type f_
    return
    Data_Type * counters_
```

#### 9.59.1 Macro Definition Documentation

#### 9.59.1.1 STATSCOUNTER\_TEMPLATE

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

#### 9.59.1.2 STATSCOUNTER\_TEMPLATE\_ARGS

```
template STATSCOUNTER_TEMPLATE_ARGS() <typename Array_Type, typename Data_Type>
```

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

#### 9.59.1.3 STATSCOUNTER\_TYPE

```
template Data_Type * STATSCOUNTER_TYPE( ) StatsCounter<Array_Type,Data_Type>
```

Definition at line 4 of file statscounter-meat.hpp.

## 9.59.2 Function Documentation

#### 9.59.2.1 clear()

```
EmptyArray clear ( )
```

## 9.59.2.2 for()

## 9.59.2.3 resize()

## 9.59.2.4 STATSCOUNTER\_TEMPLATE() [1/9]

```
STATSCOUNTER_TEMPLATE (
StatsCounter ) const
```

## 9.59.2.5 STATSCOUNTER\_TEMPLATE() [2/9]

```
STATSCOUNTER_TEMPLATE ( \sim \textit{StatsCounter} \ )
```

Definition at line 27 of file statscounter-meat.hpp.

## 9.59.2.6 STATSCOUNTER\_TEMPLATE() [3/9]

```
STATSCOUNTER_TEMPLATE (
          std::vector< std::string > ,
          get_descriptions ) const
```

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

## 9.59.2.7 STATSCOUNTER\_TEMPLATE() [4/9]

```
STATSCOUNTER_TEMPLATE (
          std::vector< std::string > ,
          get_names ) const
```

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

## 9.59.2.8 STATSCOUNTER\_TEMPLATE() [5/9]

## 9.59.2.9 STATSCOUNTER\_TEMPLATE() [6/9]

## 9.59.2.10 STATSCOUNTER\_TEMPLATE() [7/9]

```
STATSCOUNTER_TEMPLATE (

void ,

count_init )
```

## 9.59.2.11 STATSCOUNTER\_TEMPLATE() [8/9]

```
STATSCOUNTER_TEMPLATE (
void ,
reset_array ) const
```

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

## 9.59.2.12 STATSCOUNTER\_TEMPLATE() [9/9]

## 9.59.3 Variable Documentation

#### 9.59.3.1 counter

```
Data_Type& counter

Initial value:
{
    Array = counter.Array
```

Definition at line 12 of file statscounter-meat.hpp.

## 9.59.3.2 counter\_deleted

```
counter_deleted = false
```

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

#### 9.59.3.3 counters

```
counters = new Counters<Array_Type,Data_Type>((*counter.counters))
```

Definition at line 22 of file statscounter-meat.hpp.

#### 9.59.3.4 counters\_

```
Data_Type* counters_
Initial value:
{
    if (!counter_deleted)
```

delete counters

Definition at line 53 of file statscounter-meat.hpp.

## 9.59.3.5 current\_stats

```
current_stats = counter.current_stats
```

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

#### 9.59.3.6 EmptyArray

```
EmptyArray = *Array
```

Definition at line 17 of file statscounter-meat.hpp.

#### 9.59.3.7 f

```
Data_Type f_
Initial value:
{
    counters->add_counter(f_)
```

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

## 9.59.3.8 j

```
size_t j
Initial value:
{
    if (counters->size() == 0u)
```

throw std::logic\_error("No counters added: Cannot count without knowning what to count!")

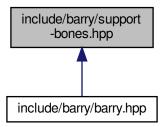
Definition at line 66 of file statscounter-meat.hpp.

#### 9.59.3.9 return

return

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

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

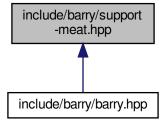


#### **Classes**

class Support < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type >
 Compute the support of sufficient statistics.

# 9.61 include/barry/support-meat.hpp File Reference

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



## **Macros**

• #define BARRY\_SUPPORT\_MEAT\_HPP 1

## 9.61.1 Macro Definition Documentation

## 9.61.1.1 BARRY\_SUPPORT\_MEAT\_HPP

#define BARRY\_SUPPORT\_MEAT\_HPP 1

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

# 9.62 include/barry/typedefs.hpp File Reference

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



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



#### **Classes**

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

# **Namespaces**

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

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

## **Typedefs**

const int EXISTS::TWO = 1
 const int EXISTS::UKNOWN = -1
 const int EXISTS::AS\_ZERO = 0
 const int EXISTS::AS\_ONE = 1

```
    typedef std::vector< std::pair< std::vector< double >, size t >> Counts type

    template<typename Cell_Type >
      using Row_type = Map< size_t, Cell< Cell_Type > >
    • template<typename Cell_Type >
      using Col_type = Map< size_t, Cell< Cell_Type > * >
    • template<typename Ta = double, typename Tb = size_t>
      using MapVec_type = std::unordered_map< std::vector< Ta >, Tb, vecHasher< Ta >>
    • template<typename Array_Type , typename Data_Type >
      using Hasher_fun_type = std::function < std::vector < double >(const Array_Type &, Data_Type *)>
          Hasher function used by the counter.

    template<typename Array_Type , typename Data_Type >

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

    std::vector < size_t > sort_array (const double *v, size_t start, size_t ncols, size_t nrows)

          Ascending sorting an array.
    • template<typename T >
      T vec_inner_prod (const T *a, const T *b, size_t n)
    • template<> double vec_inner_prod (const double *a, const double *b, size_t n)
    template<typename T >
      bool vec equal (const std::vector< T > &a, const std::vector< T > &b)
          Compares if -a- and -b- are equal.
    • template<typename T >
      bool vec_equal_approx (const std::vector< T > &a, const std::vector< T > &b, double eps=1e-100)
Variables
    • const int CHECK::BOTH = -1
    • const int CHECK::NONE = 0

    const int CHECK::ONE = 1

    const int CHECK::TWO = 2

    const int EXISTS::BOTH = -1

    const int EXISTS::NONE = 0

    • const int EXISTS::ONE = 1
```

## 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 290 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, 200
     BArray< Cell Type, Data Type >, 63
                                                       \simNode
\simBArrayCell
                                                            Node, 203
     BArrayCell< Cell_Type, Data_Type >, 74
                                                       \simPhyloRuleDynData
~BArrayCell const
                                                            PhyloRuleDynData, 212
     BArrayCell_const< Cell_Type, Data_Type >, 77
                                                       \simPowerSet
{\sim}\mathsf{BArrayDense}
                                                            PowerSet < Array_Type, Data_Rule_Type >, 215
     BArrayDense < Cell_Type, Data_Type >, 83
                                                       \simProgress
                                                            Progress, 220
\simBArrayDenseCell
     BArrayDenseCell< Cell_Type, Data_Type >, 97
                                                       \simRule
\simBArrayRow
                                                            Rule < Array_Type, Data_Type >, 222
     BArrayRow < Cell Type, Data Type >, 110
                                                       \simRules
\simBArrayRow const
                                                            Rules < Array Type, Data Type >, 225
     BArrayRow_const < Cell_Type, Data_Type >, 112
                                                       \simStatsCounter
\simBArrayVector
                                                            StatsCounter < Array_Type, Data_Type >, 230
     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 >,
                                                                 235
         118
\simCell
                                                       а
     Cell< Cell_Type >, 122
                                                            model-meat.hpp, 314
\simConstBArrayRowIter
    ConstBArrayRowlter< Cell_Type, Data_Type >,
                                                       active
                                                            Cell< Cell_Type >, 125
         127
                                                       add
\simCounter
                                                            Cell< Cell Type >, 123, 124
    Counter< Array_Type, Data_Type >, 130
                                                            FreqTable< T >, 158
\simCounters
                                                       add_array
     Counters < Array_Type, Data_Type >, 135
                                                            Model <
                                                                        Array_Type,
                                                                                         Data_Counter_Type,
\simDEFMCounterData
                                                                 Data_Rule_Type, Data_Rule_Dyn_Type >,
     Phylo rules, 43
\simDEFMData
                                                       add counter
     Phylo rules, 43
                                                            Counters < Array_Type, Data_Type >, 135, 136
\simDEFMRuleDynData
                                                            Model <
                                                                        Array Type,
                                                                                         Data Counter Type,
     Phylo rules, 43
                                                                 Data_Rule_Type, Data_Rule_Dyn_Type >,
\simEntries
                                                                 180, 181
     Entries < Cell Type >, 149
                                                            StatsCounter< Array_Type, Data_Type >, 230
\simFlock
                                                            Support<
                                                                         Array Type,
                                                                                         Data Counter Type,
     Flock, 151
                                                                 \label{eq:decomposition} Data\_Rule\_Type, \quad Data\_Rule\_Dyn\_Type \quad >,
\simFreqTable
                                                                 235
     FreqTable < T >, 158
                                                       add data
\simGeese
                                                            Flock, 151
    Geese, 164
                                                       add dims
\simModel
                                                            counters-meat.hpp, 281
    Model<
                Array Type,
                                 Data Counter Type,
         Data_Rule_Type, Data_Rule_Dyn_Type >,
                                                       add hash
                                                            Counters < Array Type, Data Type >, 136
         180
                                                       add hasher
\simNetCounterData
                                                            Model <
                                                                                         Data_Counter_Type,
                                                                        Array_Type,
    NetCounterData, 198
                                                                 Data_Rule_Type, Data_Rule_Dyn_Type >,
\simNetworkData
```

add_rule	is_dense, 66
Model < Array_Type, Data_Counter_Type,	is_empty, 66
Data_Rule_Type, Data_Rule_Dyn_Type >,	ncol, 67
181	nnozero, 67
PowerSet< Array_Type, Data_Rule_Type >, 215,	nrow, 67
216	operator*=, 67
Rules < Array_Type, Data_Type >, 226	operator(), 67
Support< Array_Type, Data_Counter_Type,	operator+=, 68
Data_Rule_Type, Data_Rule_Dyn_Type >,	operator-=, 68
235, 236	operator/=, 69
add_rule_dyn	operator=, 69
Model< Array_Type, Data_Counter_Type,	operator==, 69
Data_Rule_Type, Data_Rule_Dyn_Type >,	out_of_range, 69
181, 182	print, 69
Support< Array_Type, Data_Counter_Type,	print_n, 70
Data_Rule_Type, Data_Rule_Dyn_Type >,	reserve, 70
236	resize, 70
annotations	rm_cell, 70
Node, 204	row, 70
Array	set_data, 71
ConstBArrayRowIter< Cell_Type, Data_Type >,	swap_cells, 71
127	swap_cols, 71
array	swap_rows, 71
Node, 204	toggle_cell, 72
Phylo rules, 44	toggle_lock, 72
arrays	transpose, 72
Node, 204	visited, 73
arrays2support	zero_col, 72
Model < Array_Type, Data_Counter_Type,	zero_row, 72
Data_Rule_Type, Data_Rule_Dyn_Type >,	barray-meat-operators.hpp
191	BARRAY_TEMPLATE, 246-248
AS_ONE	BARRAY_TEMPLATE_ARGS, 247, 249
EXISTS, 57	BARRAY_TYPE, 247, 249
as_vector	COL, 247
FreqTable $<$ T $>$ , 158	for, 249
AS_ZERO	operator(), 249
EXISTS, 57	rhs, 249
at	ROW, 247
Phylo rules, 39	this, 250
PhyloCounterData, 209	barray-meat.hpp
	COL, 250
BArray	ROW, 251
BArray < Cell_Type, Data_Type >, 62, 63	BARRAY_TEMPLATE
BArray< Cell_Type, Data_Type >, 59	barray-meat-operators.hpp, 246-248
~BArray, 63	BARRAY_TEMPLATE_ARGS
BArray, 62, 63	barray-meat-operators.hpp, 247, 249
BArrayCell< Cell_Type, Data_Type >, 73	BARRAY_TYPE
BArrayCell_const< Cell_Type, Data_Type >, 73	barray-meat-operators.hpp, 247, 249
clear, 63	BArrayCell
col, 63	BArrayCell< Cell_Type, Data_Type >, 74
D, 64	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	resize, 92
operator=, 75	rm_cell, 92
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, 253, 255
BArrayDense	BDENSE_TEMPLATE_ARGS, 253, 255
BArrayDense < Cell_Type, Data_Type >, 82, 83	BDENSE_TYPE, 254, 256
BArrayDense< Cell_Type, Data_Type >, 79	COL, 254
∼BArrayDense, 83	POS, 254
BArrayDense, 82, 83	POS_N, 254
BArrayDenseCell< Cell_Type, Data_Type >, 95,	ROW, 254
99	barraydense-meat.hpp
BArrayDenseCol< Cell_Type, Data_Type >, 95,	COL, 256
102	POS, 256
BArrayDenseCol_const< Cell_Type, Data_Type >,	POS_N, 257
95	ROW, 257
BArrayDenseRow< Cell_Type, Data_Type >, 95,	ZERO_CELL, 257
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	${\sf BArrayDenseCol\_const} {< Cell\_Type, Data\_Type >},$
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, 258
operator+=, 89	barraydensecell-meat.hpp
operator-=, 90	POS, 258
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 >, 109	BArrayDenseRow_const, 107 begin, 108
BArrayDenseCol	end, 108
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,	BArrayRow< Cell_Type, Data_Type >, 110
102	BArrayRow< Cell_Type, Data_Type >, 109
BArrayDenseCell_const< Cell_Type, Data_Type	$\sim$ 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, 259	operator=, 111
POS_N, 259	operator==, 111
ZERO_CELL, 260	barrayrow-meat.hpp
BArrayDenseCol_const	BROW_TEMPLATE, 262, 263
BArrayDenseCol_const< Cell_Type, Data_Type >,	BROW_TEMPLATE_ARGS, 262
103	BROW_TYPE, 262
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	operator>, 113
BArrayDenseRow	operator>=, 113
BArrayDenseRow< Cell_Type, Data_Type >, 105	operator==, 113
BArrayDenseRow< Cell_Type, Data_Type >, 104	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	~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, 261	operator+=, 116
POS_N, 261	operator-=, 116
ZERO_CELL, 261 BArrayDenseRow_const	operator/=, 116
BArrayDenseRow_const< Cell_Type, Data_Type	operator=, 117 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< Cell_Type, Data_Type >, 93  BArrayDenseCell< Cell_Type, Data_Type >, 109	BArrayVector_const< Cell_Type, Data_Type >, 117
BArrayDenseCell_const< Cell_Type, Data_Type  BarrayDenseCell_const< Cell_Type, Data_Type	~BArrayVector_const, 118
>, 109	BArrayVector_const, 118
· , · • •	=a,

begin, 119	support-meat.hpp, 352
end, 119	BARRY_UNUSED
is_col, 119	barry-macros.hpp, 267
is_row, 119	BARRY_VERSION
operator std::vector< Cell_Type >, 119	barry.hpp, 270
operator!=, 119	BARRY_VERSION_MAYOR
operator<, 120	barry.hpp, 270
operator<=, 120	BARRY_VERSION_MINOR
operator>, 120	barry.hpp, 270
operator>=, 120	BARRY_ZERO
operator==, 120	barry-macros.hpp, 268
size, 121	BARRY_ZERO_DENSE
barry, 55	barry-macros.hpp, 268
barry-configuration.hpp	BARRY_ZERO_NETWORK
BARRY_CHECK_SUPPORT, 265	network.hpp, 296
BARRY_ISFINITE, 265	BARRY_ZERO_NETWORK_DENSE
BARRY_MAX_NUM_ELEMENTS, 265	network.hpp, 296
BARRY_SAFE_EXP, 265	BDENSE_TEMPLATE
Map, 266	barraydense-meat-operators.hpp, 253, 255
printf_barry, 265	BDENSE_TEMPLATE_ARGS
barry-debug.hpp	barraydense-meat-operators.hpp, 253, 255
BARRY_DEBUG_LEVEL, 266	BDENSE_TYPE
barry-macros.hpp	barraydense-meat-operators.hpp, 254, 256
BARRY_ONE, 267	begin
BARRY_ONE_DENSE, 267	BArrayDenseCol< Cell_Type, Data_Type >, 101
BARRY_UNUSED, 267	BArrayDenseCol_const< Cell_Type, Data_Type >,
BARRY_ZERO, 268	103
BARRY_ZERO_DENSE, 268	BArrayDenseRow< Cell_Type, Data_Type >, 105
barry.hpp	BArrayDenseRow_const< Cell_Type, Data_Type
BARRY_HPP, 269	>, 108
BARRY_VERSION, 270	BArrayVector< Cell_Type, Data_Type >, 115
BARRY_VERSION_MAYOR, 270	BArrayVector_const $<$ Cell_Type, Data_Type $>$ ,
BARRY_VERSION_MINOR, 270	119
COUNTER_FUNCTION, 270	PhyloCounterData, 209
COUNTER_LAMBDA, 270	PowerSet < Array_Type, Data_Rule_Type >, 216
RULE_FUNCTION, 271	Rules< Array_Type, Data_Type >, 226
RULE_LAMBDA, 271	blengths
barry::counters, 55	NodeData, 207
barry::counters::network, 56	ВОТН
BARRY_CHECK_SUPPORT	CHECK, 56
barry-configuration.hpp, 265	EXISTS, 57
BARRY_DEBUG_LEVEL	BROW_TEMPLATE
barry-debug.hpp, 266	barrayrow-meat.hpp, 262, 263
BARRY_HPP	BROW_TEMPLATE_ARGS
barry.hpp, 269	barrayrow-meat.hpp, 262
BARRY_ISFINITE	BROW_TYPE
barry-configuration.hpp, 265	barrayrow-meat.hpp, 262
BARRY_MAX_NUM_ELEMENTS	aala
barry-configuration.hpp, 265	calc
BARRY_ONE	PowerSet < Array_Type, Data_Rule_Type >, 216
barry-macros.hpp, 267	Support< Array_Type, Data_Counter_Type,
BARRY_ONE_DENSE	Data_Rule_Type, Data_Rule_Dyn_Type >,
barry-macros.hpp, 267	236
BARRY_PROGRESS_BAR_WIDTH	calc_reduced_sequence
progress.hpp, 344	Geese, 164
BARRY_SAFE_EXP	calc_sequence
barry-configuration.hpp, 265	Geese, 165
BARRY_SUPPORT_MEAT_HPP	Cell Cell Type > 100 100
	Cell< Cell_Type >, 122, 123

Cell< Cell_Type >, 121	coordiantes_n_free
∼Cell, 122	Support< Array_Type, Data_Counter_Type,
active, 125	Data_Rule_Type, Data_Rule_Dyn_Type >,
add, 123, 124	240
Cell, 122, 123	coordiantes_n_locked
operator Cell_Type, 124	Support< Array_Type, Data_Counter_Type,
operator!=, 124	Data_Rule_Type, Data_Rule_Dyn_Type >,
operator=, 124, 125	241
operator==, 125	coordinates_free
value, 125	PowerSet < Array_Type, Data_Rule_Type >, 218
visited, 125	Support< Array_Type, Data_Counter_Type,
Cell_const< Cell_Type >, 126	Data_Rule_Type, Data_Rule_Dyn_Type >,
change_stats	241
Support< Array_Type, Data_Counter_Type,	
Data_Rule_Type, Data_Rule_Dyn_Type >,	PowerSet < Array_Type, Data_Rule_Type >, 218
240	Support< Array_Type, Data_Counter_Type,
CHECK, 56	Data_Rule_Type, Data_Rule_Dyn_Type >,
BOTH, 56	241
NONE, 56	count
ONE, 56	Counter< Array_Type, Data_Type >, 131
TWO, 56	count_all
clear	StatsCounter< Array_Type, Data_Type >, 230
BArray< Cell_Type, Data_Type >, 63	count_current
BArrayDense < Cell_Type, Data_Type >, 83	StatsCounter< Array_Type, Data_Type >, 231
FreqTable $<$ T $>$ , 158	count_fun
statscounter-meat.hpp, 347	Counter< Array_Type, Data_Type >, 132
COL	counters-meat.hpp, 276
barray-meat-operators.hpp, 247	count_fun_
barray-meat.hpp, 250	counters-meat.hpp, 281
barraydense-meat-operators.hpp, 254	model-meat.hpp, 314
barraydense-meat.hpp, 256	count_init
col	StatsCounter< Array_Type, Data_Type >, 231
BArray< Cell_Type, Data_Type >, 63	Counter
BArrayDense< Cell_Type, Data_Type >, 84	Counter< Array_Type, Data_Type >, 130
Col_type	counter
typedefs.hpp, 355	counters-meat.hpp, 282
colnames	model-meat.hpp, 314
Flock, 152	statscounter-meat.hpp, 349
Geese, 165	Counter< Array_Type, Data_Type >, 128
Model< Array_Type, Data_Counter_Type,	∼Counter, 130
Data_Rule_Type, Data_Rule_Dyn_Type >,	count, 131
182	count_fun, 132
colsum	Counter, 130
BArrayDense < Cell_Type, Data_Type >, 84	data, 132
conditional_prob	desc, 133
Model< Array_Type, Data_Counter_Type,	get_description, 131
Data_Rule_Type, Data_Rule_Dyn_Type >,	get_hasher, 131
182	get_name, 131
ConstBArrayRowlter	hasher_fun, 133
ConstBArrayRowlter< Cell_Type, Data_Type >,	init, 131
127	init_fun, 133
ConstBArrayRowlter< Cell_Type, Data_Type >, 126	name, 133
~ConstBArrayRowlter, 127	operator=, 131, 132
Array, 127	set_hasher, 132
ConstBArrayRowlter, 127	counter_
current_col, 128	counters-meat.hpp, 282
current_row, 128	counter_absdiff
iter, 128	Network counters, 26

counter_co_opt	counter_gains
Counting, 17	Counting, 18
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, 289	Counting, 19
counter_css_census02	Phylo counters, 49
network-css.hpp, 289	counter genes changing
counter_css_census03	Counting, 19
network-css.hpp, 289	Phylo counters, 50
counter_css_census04	counter_idegree
network-css.hpp, 290	Network counters, 29
counter_css_census05	counter_idegree15
network-css.hpp, 290	Network counters, 29
counter_css_census06	counter isolates
network-css.hpp, 290	Network counters, 29, 30
counter_css_census07	counter_istar2
network-css.hpp, 290	Network counters, 30
counter_css_census08	counter_k_genes_changing
network-css.hpp, 291	Counting, 19
counter_css_census09	Phylo counters, 50
network-css.hpp, 291	COUNTER LAMBDA
counter_css_census10	barry.hpp, 270
network-css.hpp, 291	counter_less_than_p_prop_genes_changing
counter_css_completely_false_recip_comiss	Counting, 19
network-css.hpp, 291	Phylo counters, 50
counter_css_completely_false_recip_omiss	counter_logit_intercept
network-css.hpp, 292	Network counters, 30
counter_css_mixed_recip	counter_longest
network-css.hpp, 292	Counting, 20
counter_css_partially_false_recip_commi	Phylo counters, 50
network-css.hpp, 292	counter_loss
counter_css_partially_false_recip_omiss	Counting, 20
network-css.hpp, 293	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, 350	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
191	counter_nodeocov
Counter_fun_type	Network counters, 31
typedefs.hpp, 355	counter_odegree
COUNTER_FUNCTION	Network counters, 31, 32
barry.hpp, 270	counter_odegree15

Network counters, 32	$\sim$ Counters, 135
counter_ones	add_counter, 135, 136
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, 281
Counting, 21	count_fun, 276
Phylo counters, 52	count_fun_, 281
counter_overall_loss	counter, 282
Counting, 22	counter_, 282
Phylo counters, 52	COUNTER_TEMPLATE, 274, 276, 277
counter_pairwise_first_gain	COUNTER TEMPLATE ARGS, 274
Counting, 22	COUNTER TYPE, 275
Phylo counters, 53	COUNTERS TEMPLATE, 275, 277–279
counter_pairwise_neofun_singlefun	COUNTERS TEMPLATE ARGS, 275
Counting, 22	COUNTERS_TYPE, 275
Phylo counters, 53	data, 279
counter_pairwise_overall_change	data_, 282
Counting, 22	desc, 279
Phylo counters, 53	desc_, 282
counter_pairwise_preserving	for, 279
Counting, 23	fun, 283
Phylo counters, 53	fun_, 283
counter_preserve_pseudogene	hasher, 279, 280
Counting, 23	hasher_fun, 280
Phylo counters, 54	hasher_fun_, 283
counter_prop_genes_changing	i, 283
Counting, 23	if, 280
Phylo counters, 54	
-	init_fun, 281
counter_subfun	init_fun_, 284
Counting, 23	j, 284
Phylo counters, 54	name, 281
COUNTER_TEMPLATE	name_, 284
counters-meat.hpp, 274, 276, 277	noexcept, 284
COUNTER_TEMPLATE_ARGS	res, 284
counters-meat.hpp, 274	return, 285
counter_transition	TMP_HASHER_CALL, 275
Network counters, 33	counters.hpp
counter_transition_formula	DEFM_COUNTER, 322
Network counters, 34	DEFM_COUNTER_LAMBDA, 322
counter_ttriads	DEFM_RULE, 323
Network counters, 34	DEFM_RULE_LAMBDA, 323
COUNTER_TYPE	DEFM_RULEDYN_LAMBDA, 323
counters-meat.hpp, 275	UNI_SUB, 323
Counters	counters_
Counters < Array_Type, Data_Type >, 134, 135	model-meat.hpp, 314
counters	statscounter-meat.hpp, 350
Model< Array_Type, Data_Counter_Type,	COUNTERS_TEMPLATE
Data_Rule_Type, Data_Rule_Dyn_Type >,	counters-meat.hpp, 275, 277–279
191	COUNTERS_TEMPLATE_ARGS
statscounter-meat.hpp, 350	counters-meat.hpp, 275
Counters < Array_Type, Data_Type >, 134	COUNTERS_TYPE

counters-meat.hpp, 275	CSS_CHECK_SIZE_INIT
Counting, 13	network-css.hpp, 287
counter_co_opt, 17	CSS_MATCH_TYPE
counter_cogain, 18	network-css.hpp, 288
counter_gains, 18	CSS_NET_COUNTER_LAMBDA_INIT
counter_gains_from_0, 18	network-css.hpp, 288
counter_gains_k_offspring, 19	CSS_PERCEIVED_CELLS
counter_genes_changing, 19	network-css.hpp, 288
counter_k_genes_changing, 19	CSS_SIZE
counter_less_than_p_prop_genes_changing, 19	network-css.hpp, 288
counter_longest, 20	CSS_TRUE_CELLS
counter_loss, 20	network-css.hpp, 289
counter_maxfuns, 20	cumprob
counter_neofun, 20	model-meat.hpp, 315
counter_neofun_a2b, 21	current_col
counter_overall_changes, 21	ConstBArrayRowIter< Cell_Type, Data_Type >,
counter_overall_gains, 21	128
counter_overall_gains_from_0, 21	current_row
counter_overall_loss, 22	ConstBArrayRowIter< Cell_Type, Data_Type >,
counter_pairwise_first_gain, 22	128
counter_pairwise_neofun_singlefun, 22	current_stats
counter_pairwise_overall_change, 22	statscounter-meat.hpp, 350
counter_pairwise_preserving, 23	Support< Array_Type, Data_Counter_Type,
counter_preserve_pseudogene, 23	Data_Rule_Type, Data_Rule_Dyn_Type >,
counter_prop_genes_changing, 23	241
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 >, 223
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, 279
Counts  Phylograph 44	PowerSet< Array Type, Data Rule Type >, 218
Phylo rules, 44	data_
PhyloRuleDynData, 212 Counts type	counters-meat.hpp, 282
typedefs.hpp, 355	model-meat.hpp, 315
covar sort	Data Counter Type
Phylo rules, 44	model-meat.hpp, 315
covar_used	Data_Rule_Type
Phylo rules, 44	model-meat.hpp, 315
covariates	default_val
Phylo rules, 44	BArray< Cell_Type, Data_Type >, 64
CSS APPEND	BArrayDense< Cell_Type, Data_Type >, 85
network-css.hpp, 286	DEFM, 139
CSS_CASE_ELSE	DEFM, 140
network-css.hpp, 287	get_column_major, 140
CSS_CASE_PERCEIVED	get_ID, 140
network-css.hpp, 287	get_m_order, 140
CSS_CASE_TRUTH	get_model, 141
network-css.hpp, 287	get_n_covars, 141
CSS_CHECK_SIZE	get_n_obs, 141
network-css.hpp, 287	get_n_rows, 141
·	get_n_y, 141

get_X, 141 get_X_names, 142 get_Y, 142	DEFMStatsCounter Phylo rules, 38 DEFMSupport
get_Y_names, 142	Phylo rules, 38
init, 142	delete_counters
is_motif, 142	Model< Array_Type, Data_Counter_Type,
logodds, 142	Data_Rule_Type, Data_Rule_Dyn_Type >,
motif_census, 143	192
print, 143	Support< Array_Type, Data_Counter_Type,
set_names, 143	Data_Rule_Type, Data_Rule_Dyn_Type >,
simulate, 143	241
defm, 57	delete_rengine
defm-meat.hpp	Geese, 172
DEFM_LOOP_ARRAYS, 327	Model < Array_Type, Data_Counter_Type,
DEFM_RANGES, 327	Data_Rule_Type, Data_Rule_Dyn_Type >,
keygen_defm, 327	192
defm-types.hpp	delete_rules
DEFMArray, 328	Model < Array_Type, Data_Counter_Type,
DEFM_COUNTER	Data_Rule_Type, Data_Rule_Dyn_Type >,
counters.hpp, 322	192
DEFM_COUNTER_LAMBDA	model-meat.hpp, 315
counters.hpp, 322	Support
DEFM_LOOP_ARRAYS	Data_Rule_Type, Data_Rule_Dyn_Type >, 242
defm-meat.hpp, 327	
defm_motif_parser	delete_rules_dyn  Model< Array Type. Data Counter Type.
formula.hpp, 329 DEFM_RANGES	Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,
defm-meat.hpp, 327	192
DEFM_RULE	model-meat.hpp, 316
counters.hpp, 323	Support< Array_Type, Data_Counter_Type,
DEFM_RULE_LAMBDA	Data_Rule_Type, Data_Rule_Dyn_Type >,
counters.hpp, 323	242
DEFM_RULEDYN_LAMBDA	delete_support
counters.hpp, 323	Geese, 172
DEFMArray	desc
defm-types.hpp, 328	Counter< Array_Type, Data_Type >, 133
DEFMCounter	counters-meat.hpp, 279
Phylo rules, 37	desc_
DEFMCounterData, 144	counters-meat.hpp, 282
Phylo rules, 39	directed
DEFMCounters	NetworkData, 200
Phylo rules, 37	duplication
DEFMData, 144	Node, 205
Phylo rules, 39	NodeData, 208
DEFMModel	PhyloRuleDynData, 212
Phylo rules, 37	
DEFMRule	else
Phylo rules, 38	model-meat.hpp, 316
DEFMRuleData, 146	empty
Phylo rules, 40	PhyloCounterData, 209
DEFMRuleDyn	EmptyArray
Phylo rules, 38	PowerSet < Array_Type, Data_Rule_Type >, 218
DEFMRuleDynData, 147	statscounter-meat.hpp, 350
Phylo rules, 40	end  PArroy Donos Col. Coll Type Data Type > 101
DEFMRules	BArrayDenseCol const < Coll Type, Data_Type >, 101
Phylorules, 38	BArrayDenseCol_const< Cell_Type, Data_Type >, 103
DEFMRulesDyn	
Phylo rules, 38	BArrayDenseRow< Cell_Type, Data_Type >, 105

BArrayDenseRow_const< Cell_Type, Data_Type	model, 156
>, 108	nfunctions, 156
BArrayVector< Cell_Type, Data_Type >, 115	nfuns, 153
BArrayVector_const< Cell_Type, Data_Type >,	nleafs, 154
119	nnodes, 154
PhyloCounterData, 210	nterms, 154
PowerSet < Array_Type, Data_Rule_Type >, 216	ntrees, 154
Progress, 221	operator(), 154
Rules < Array_Type, Data_Type >, 226	parse_polytomies, 155
Entries	print, 155
Entries< Cell_Type >, 148	rengine, 156
Entries < Cell_Type >, 148	set_seed, 155
~Entries, 149	support_size, 155
Entries, 148	flush_data
resize, 149	BArray< Cell_Type, Data_Type >, 64
source, 149	for
target, 149	barray-meat-operators.hpp, 249
val, 149	counters-meat.hpp, 279
•	• • •
etype_default	model-meat.hpp, 304
Geese, 172	statscounter-meat.hpp, 347
etype_duplication	force_new
Geese, 172	model-meat.hpp, 316
etype_either	formula.hpp
Geese, 172	defm_motif_parser, 329
etype_speciation	FreqTable
Geese, 173	FreqTable < T >, 158
eval_rules_dyn	FreqTable < T >, 157
Support< Array_Type, Data_Counter_Type,	∼FreqTable, 158
Data_Rule_Type, Data_Rule_Dyn_Type >,	add, 158
237	as_vector, 158
EXISTS, 57	clear, 158
AS_ONE, 57	FreqTable, 158
AS_ZERO, 57	get_data, 159
BOTH, 57	get_index, 159
NONE, 58	make_hash, 159
ONE, 58	print, 159
TWO, 58	reserve, 159
UKNOWN, 58	size, 160
	fun
f_	counters-meat.hpp, 283
statscounter-meat.hpp, 351	fun_
first_calc_done	counters-meat.hpp, 283
Model< Array_Type, Data_Counter_Type,	model-meat.hpp, 316
Data_Rule_Type, Data_Rule_Dyn_Type >,	• • •
192	Geese, 160
Flock, 150	$\sim$ Geese, 164
$\sim$ Flock, 151	calc_reduced_sequence, 164
add_data, 151	calc_sequence, 165
colnames, 152	colnames, 165
dat, 156	delete_rengine, 172
Flock, 151	delete_support, 172
get_counters, 152	etype_default, 172
get_model, 152	etype_duplication, 172
get_stats_support, 152	etype_either, 172
get_stats_target, 152	etype_speciation, 173
get_support_fun, 153	Geese, 163, 164
init, 153	get_annotated_nodes, 165
initialized, 156	get_annotations, 165
likelihood_joint, 153	get_counters, 165
<b>_</b> ,	5,

get model 166	Countary / Array Time Data Time > 126
get_model, 166 get_probabilities, 166	Counters < Array_Type, Data_Type >, 136
<del>-</del>	gen_key  Model< Array Type. Data Counter Type.
get_rengine, 166	3= 31 / = = 31 /
get_states, 166	Data_Rule_Type, Data_Rule_Dyn_Type >
get_support_fun, 166	183
inherit_support, 167	get_annotated_nodes
init, 167	Geese, 165
init_node, 167	get_annotations
initialized, 173	Geese, 165
likelihood, 167	get_arrays2support
likelihood exhaust, 167	Model< Array_Type, Data_Counter_Type,
map_to_state_id, 173	Data_Rule_Type, Data_Rule_Dyn_Type >
nannotations, 168	183
nfunctions, 173	get cell
nfuns, 168	BArray< Cell_Type, Data_Type >, 64
nleafs, 168	BArrayDense< Cell_Type, Data_Type >, 85
nnodes, 168	
	get_col_vec
nodes, 173	BArray Cell_Type, Data_Type >, 65
nterms, 168	BArrayDense < Cell_Type, Data_Type >, 85, 86
observed_counts, 169	get_column_major
operator=, 169	DEFM, 140
parse_polytomies, 169	get_counters
predict, 169	Flock, 152
predict_backend, 170	Geese, 165
predict_exhaust, 170	Model < Array_Type, Data_Counter_Type,
predict_exhaust_backend, 170	Data_Rule_Type, Data_Rule_Dyn_Type >
predict_sim, 170	183
print, 170	PhyloCounterData, 210
print_nodes, 171	StatsCounter< Array_Type, Data_Type >, 231
print_observed_counts, 171	Support< Array_Type, Data_Counter_Type,
pset_loc, 173	Data_Rule_Type, Data_Rule_Dyn_Type >:
reduced_sequence, 174	237
sequence, 174	get_counts
set_seed, 171	Support< Array_Type, Data_Counter_Type
simulate, 171	Data_Rule_Type, Data_Rule_Dyn_Type >
support_size, 171	237
update_annotations, 171	get_current_stats
geese, 58	Support< Array_Type, Data_Counter_Type
geese-bones.hpp	Data_Rule_Type, Data_Rule_Dyn_Type >
INITIALIZED, 333	237
keygen_full, 333	get_data
RULE FUNCTION, 333	BArrayDense < Cell_Type, Data_Type >, 86
vec_diff, 333	FreqTable < T >, 159
vector_caster, 334	PowerSet < Array_Type, Data_Rule_Type >, 216
geese-types.hpp	Support< Array_Type, Data_Counter_Type,
PhyloArray, 340	Data_Rule_Type, Data_Rule_Dyn_Type >:
PhyloCounter, 340	238
PhyloCounters, 340	get_data_ptr
PhyloModel, 340	PowerSet < Array_Type, Data_Rule_Type >, 217
PhyloPowerSet, 340	get_description
PhyloRule, 341	Counter< Array_Type, Data_Type >, 131
PhyloRuleData, 341	Rule < Array_Type, Data_Type >, 223
PhyloRuleDyn, 341	get_descriptions
PhyloRules, 341	Counters < Array_Type, Data_Type >, 136
PhyloRulesDyn, 341	Rules < Array_Type, Data_Type >, 226
PhyloStatsCounter, 341	StatsCounter< Array_Type, Data_Type >, 231
PhyloSupport, 342	get_entries
gen_hash	BArray< Cell_Type, Data_Type >, 65
9	··· · · · · · · · · · · · · · · · ·

BArrayDense < Cell_Type, Data_Type >, 86	184
get_hasher	get_row_vec
Counter< Array_Type, Data_Type >, 131	BArray< Cell_Type, Data_Type >, 65
get_ID	BArrayDense < Cell_Type, Data_Type >, 86, 87
DEFM, 140	get_rules
get_index	Model < Array_Type, Data_Counter_Type,
FreqTable $<$ T $>$ , 159	Data_Rule_Type, Data_Rule_Dyn_Type >,
get_last_name	185
Counting, 24	Support< Array_Type, Data_Counter_Type,
get_m_order	Data_Rule_Type, Data_Rule_Dyn_Type >,
DEFM, 140	238
get_model	get rules dyn
DEFM, 141	Model< Array_Type, Data_Counter_Type,
Flock, 152	Data_Rule_Type, Data_Rule_Dyn_Type >,
Geese, 166	185
get_n_covars	Support< Array_Type, Data_Counter_Type,
DEFM, 141	Data_Rule_Type, Data_Rule_Dyn_Type >,
get_n_obs	238
DEFM, 141	get seq
get_n_rows	Rules < Array_Type, Data_Type >, 227
DEFM, 141	get states
get_n_y	Geese, 166
DEFM, 141	get_stats_support
get name	Flock, 152
Counter< Array_Type, Data_Type >, 131	Model < Array_Type, Data_Counter_Type,
Rule < Array_Type, Data_Type >, 223	Data_Rule_Type, Data_Rule_Dyn_Type >,
get names	185
Counters< Array_Type, Data_Type >, 137	get_stats_target
Rules < Array_Type, Data_Type >, 227	Flock, 152
StatsCounter< Array_Type, Data_Type >, 231	Model < Array_Type, Data_Counter_Type,
get_norm_const	Data_Rule_Type, Data_Rule_Dyn_Type >,
Model< Array_Type, Data_Counter_Type,	185
Data_Rule_Type, Data_Rule_Dyn_Type >,	get_support_fun
183	Flock, 153
get_parent	Geese, 166
Node, 203	Model < Array_Type, Data_Counter_Type,
get_probabilities	Data_Rule_Type, Data_Rule_Dyn_Type >,
Geese, 166	185
get_pset	get_X
Model< Array_Type, Data_Counter_Type,	DEFM, 141
Data_Rule_Type, Data_Rule_Dyn_Type >,	get_X_names
183	DEFM, 142
get_pset_arrays	get_Y
Model< Array_Type, Data_Counter_Type,	DEFM, 142
Data_Rule_Type, Data_Rule_Dyn_Type >,	get_Y_names
184	DEFM, 142
get_pset_probs	
Model< Array_Type, Data_Counter_Type,	hasher
Data_Rule_Type, Data_Rule_Dyn_Type >,	counters-meat.hpp, 279, 280
184	hasher_fun
get_pset_stats	Counter< Array_Type, Data_Type >, 133
Model< Array_Type, Data_Counter_Type,	counters-meat.hpp, 280
Data_Rule_Type, Data_Rule_Dyn_Type >,	hasher_fun_
184	counters-meat.hpp, 283
get_rengine	Hasher_fun_type
Geese, 166	typedefs.hpp, 355
Model< Array_Type, Data_Counter_Type,	hashes
Data_Rule_Type, Data_Rule_Dyn_Type >,	Support< Array_Type, Data_Counter_Type,
	Data_Rule_Type, Data_Rule_Dyn_Type >,

242	include/barry/models/geese.hpp, 331
hashes_initialized	include/barry/models/geese/counters.hpp, 324
Support< Array_Type, Data_Counter_Type,	include/barry/models/geese/flock-bones.hpp, 331
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/models/geese/flock-meat.hpp, 332
242	include/barry/models/geese/geese-bones.hpp, 332
242	include/barry/models/geese/geese-meat-constructors.hpp,
i	
counters-meat.hpp, 283	334
model-meat.hpp, 316	include/barry/models/geese/geese-meat-likelihood.hpp,
	335
i_matches	include/barry/models/geese/geese-meat-likelihood_exhaust.hpp,
model-meat.hpp, 317	336
id	include/barry/models/geese/geese-meat-predict.hpp,
Node, 205	336
idx	include/barry/models/geese/geese-meat-predict_exhaust.hpp,
Phylo rules, 40, 41	337
if	include/barry/models/geese/geese-meat-predict_sim.hpp,
counters-meat.hpp, 280	337
model-meat.hpp, 304, 305	
IF_MATCHES	include/barry/models/geese/geese-meat-simulate.hpp,
Counting, 15	338
IF_NOTMATCHES	include/barry/models/geese/geese-meat.hpp, 338
	include/barry/models/geese/geese-node-bones.hpp,
Counting, 15	339
include/barry/barray-bones.hpp, 245	include/barry/models/geese/geese-types.hpp, 339
include/barry/barray-iterator.hpp, 245	include/barry/powerset-bones.hpp, 342
include/barry/barray-meat-operators.hpp, 246	include/barry/powerset-meat.hpp, 343
include/barry/barray-meat.hpp, 250	include/barry/progress.hpp, 343
include/barry/barraycell-bones.hpp, 251	include/barry/rules-bones.hpp, 344
include/barry/barraycell-meat.hpp, 252	include/barry/rules-meat.hpp, 345
include/barry/barraydense-bones.hpp, 252	include/barry/statscounter-bones.hpp, 345
include/barry/barraydense-meat-operators.hpp, 253	•
include/barry/barraydense-meat.hpp, 256	include/barry/statscounter-meat.hpp, 346
include/barry/barraydensecell-bones.hpp, 257	include/barry/support-bones.hpp, 351
include/barry/barraydensecell-meat.hpp, 258	include/barry/support-meat.hpp, 352
• • • • • • • • • • • • • • • • • • • •	include/barry/typedefs.hpp, 353
include/barry/barraydensecol-bones.hpp, 259	indices
include/barry/barraydenserow-bones.hpp, 260	NetCounterData, 198
include/barry/barrayrow-bones.hpp, 261	Phylo rules, 44, 45
include/barry/barrayrow-meat.hpp, 261	inherit_support
include/barry/barrayvector-bones.hpp, 263	Geese, 167
include/barry/barrayvector-meat.hpp, 264	init
include/barry/barry-configuration.hpp, 264	Counter< Array_Type, Data_Type >, 131
include/barry/barry-debug.hpp, 266	DEFM, 142
include/barry/barry-macros.hpp, 267	Flock, 153
include/barry/barry.hpp, 268	Geese, 167
include/barry/cell-bones.hpp, 271	
include/barry/cell-meat.hpp, 272	Phylo rules, 45
include/barry/col-bones.hpp, 272	init_fun
· · · · · · · · · · · · · · · · · · ·	Counter< Array_Type, Data_Type >, 133
include/barry/counters-bones.hpp, 272	counters-meat.hpp, 281
include/barry/counters-meat.hpp, 273	init_fun_
include/barry/counters/network-css.hpp, 285	counters-meat.hpp, 284
include/barry/counters/network.hpp, 293	model-meat.hpp, 317
include/barry/freqtable.hpp, 300	init_node
include/barry/model-bones.hpp, 301	Geese, 167
include/barry/model-meat.hpp, 301	init_support
include/barry/models/defm.hpp, 320	PowerSet < Array_Type, Data_Rule_Type >, 217
include/barry/models/defm/counters.hpp, 321	
include/barry/models/defm/defm-bones.hpp, 326	Support< Array_Type, Data_Counter_Type,
include/barry/models/defm/defm-meat.hpp, 326	Data_Rule_Type, Data_Rule_Dyn_Type >,
- · · · · · · · · · · · · · · · · · · ·	238
include/barry/models/defm/defm-types.hpp, 328	INITIALIZED
include/barry/models/defm/formula.hpp, 329	

geese-bones.hpp, 333	likelihood
initialized	Geese, 167
Flock, 156	Model< Array_Type, Data_Counter_Type,
Geese, 173	${\sf Data\_Rule\_Type,\ \ Data\_Rule\_Dyn\_Type\ >,}$
insert_cell	186
BArray< Cell_Type, Data_Type >, 66	likelihood_
BArrayDense< Cell_Type, Data_Type >, 87	model-meat.hpp, 305
model-meat.hpp, 305	likelihood_exhaust
is_col	Geese, 167
BArrayVector< Cell_Type, Data_Type >, 115	likelihood_joint
BArrayVector_const< Cell_Type, Data_Type >,	Flock, 153
119	likelihood_total
is_dense	Model< Array_Type, Data_Counter_Type,
BArray< Cell_Type, Data_Type >, 66	Data_Rule_Type, Data_Rule_Dyn_Type >,
BArrayDense< Cell_Type, Data_Type >, 87	187
IS_DUPLICATION	locator
Counting, 15	model-meat.hpp, 317
IS_EITHER	logical
Counting, 15	Phylo rules, 45
is_empty	logodds
BArray< Cell_Type, Data_Type >, 66	DEFM, 142
BArrayDense< Cell_Type, Data_Type >, 88	•
is_leaf	M
Node, 204	PowerSet < Array_Type, Data_Rule_Type >, 218
is_motif	Support< Array_Type, Data_Counter_Type,
DEFM, 142	Data_Rule_Type, Data_Rule_Dyn_Type >,
Phylo rules, 45	242
is_row	MAKE_DEFM_HASHER
BArrayVector< Cell_Type, Data_Type >, 115	Counting, 16
BArrayVector_const< Cell_Type, Data_Type >,	MAKE_DUPL_VARS
119	Counting, 16
IS_SPECIATION	make_hash
Counting, 16	FreqTable $<$ T $>$ , 159
is_true	Мар
Phylo rules, 41	barry-configuration.hpp, 266
iter	map_to_state_id
ConstBArrayRowIter< Cell_Type, Data_Type >,	Geese, 173
128	MapVec_type
	typedefs.hpp, 356
j	max_num_elements
counters-meat.hpp, 284	Support< Array_Type, Data_Counter_Type,
model-meat.hpp, 317	$Data_Rule_Type, Data_Rule_Dyn_Type >,$
statscounter-meat.hpp, 351	243
	Model
k	Model < Array_Type, Data_Counter_Type,
model-meat.hpp, 317	Data_Rule_Type, Data_Rule_Dyn_Type >,
key	179
model-meat.hpp, 317	model
keygen_defm	Flock, 156
defm-meat.hpp, 327	Model < Array_Type, Data_Counter_Type, Data_Rule_Type
keygen_full	Data_Rule_Dyn_Type >, 175
geese-bones.hpp, 333	$\sim$ Model, 180
keys2support	add_array, 180
Model< Array_Type, Data_Counter_Type,	add_counter, 180, 181
Data_Rule_Type, Data_Rule_Dyn_Type >,	add_hasher, 181
193	add_rule, 181
	add_rule_dyn, 181, 182
lb	arrays2support, 191
PhyloRuleDynData, 213	colnames, 182
	,

conditional_prob, 182	with_pset, 197
counter_fun, 191	model-meat.hpp
counters, 191	a, 314
delete_counters, 192	count_fun_, 314
delete_rengine, 192	counter, 314
delete_rules, 192	counters_, 314
delete_rules_dyn, 192	cumprob, 315
first_calc_done, 192	data_, 315
gen_key, 183	Data_Counter_Type, 315
get_arrays2support, 183	Data_Rule_Type, 315
get_counters, 183	delete_rules, 315
get_norm_const, 183	delete_rules_dyn, 316
get_pset, 183	else, 316
get_pset_arrays, 184	for, 304
get_pset_probs, 184	force_new, 316
get_pset_stats, 184	fun_, 316
get_rengine, 184	i, 316
get_rules, 185	i_matches, 317
get_rules_dyn, 185	if, 304, 305
get_stats_support, 185	init_fun_, 317
get_stats_target, 185	insert_cell, 305
get_support_fun, 185	j, 317
keys2support, 193	k, 317
likelihood, 186	key, 317
likelihood_total, 187	likelihood_, 305
Model, 179	locator, 317
normalizing_constants, 193	MODEL_TEMPLATE, 303, 306-312
nrules, 187	MODEL_TEMPLATE_ARGS, 304
nrules_dyn, 187	MODEL_TYPE, 304
nterms, 187	params, 318
operator=, 187	probs, 318
params_last, 193	pset_arrays, 318
print, 188	push_back, 312
print_stats, 188	r, 318
pset_arrays, 193	reserve, 312
pset_probs, 194	return, 313, 318
pset_stats, 194	rule_fun_, 318
rengine, 194	rules, 319
rules, 194	rules_, 319
rules_dyn, 195	rules_dyn, 319
sample, 188	set_counters, 313
set_counters, 189	set_rules, 313
set_rengine, 189	set_rules_dyn, 313
set_rules, 189	size, 313
set_rules_dyn, 189	stats, 319
set_seed, 189	stats_support_n_arrays, 319
set_transform_model, 190	temp_stats, 320
size, 190	tmp_counts, 320
size_unique, 190	update_normalizing_constant, 313
stats_support, 195	urand, 314
stats_support_n_arrays, 195	MODEL_TEMPLATE
stats_target, 195	model-meat.hpp, 303, 306-312
store_psets, 190	MODEL_TEMPLATE_ARGS
support_fun, 196	model-meat.hpp, 304
support_size, 191	MODEL TYPE
transform_model, 191	model-meat.hpp, 304
transform_model_fun, 196	motif_census
transform_model_term_names, 196	DEFM, 143
	, -

N	counter_fixed_effect, 28
PowerSet< Array_Type, Data_Rule_Type >, 219	counter idegree, 29
Support< Array_Type, Data_Counter_Type,	counter_idegree15, 29
Data_Rule_Type, Data_Rule_Dyn_Type >,	counter_isolates, 29, 30
243	counter_istar2, 30
n_counters	counter_logit_intercept, 30
Support< Array_Type, Data_Counter_Type,	counter_mutual, 30
Data_Rule_Type, Data_Rule_Dyn_Type >,	counter_nodecov, 31
243	counter_nodeicov, 31
n_free	counter_nodematch, 31
PowerSet < Array_Type, Data_Rule_Type >, 219	counter_nodeocov, 31
n_locked	counter_odegree, 31, 32
PowerSet < Array_Type, Data_Rule_Type >, 219	counter_odegree15, 32
name	counter_ones, 32
Counter< Array_Type, Data_Type >, 133	counter_ostar2, 33
counters-meat.hpp, 281	counter_transition, 33
name_	counter_transition_formula, 34
counters-meat.hpp, 284	counter_ttriads, 34
nannotations	NETWORK_COUNTER, 34
Geese, 168	rules_dont_become_zero, 34
narray	rules_markov_fixed, 35
Node, 205	network-css.hpp
ncol	counter_css_census01, 289
BArray< Cell_Type, Data_Type >, 67	counter_css_census02, 289
BArrayDense< Cell_Type, Data_Type >, 88	counter_css_census03, 289
Phylo rules, 41	counter_css_census04, 290
NET_C_DATA_IDX	counter_css_census05, 290
network.hpp, 297	counter_css_census06, 290
NET_C_DATA_NUM	counter_css_census07, 290
network.hpp, 297	counter_css_census08, 291
NetCounter	counter_css_census09, 291
network.hpp, 298	counter_css_census10, 291
NetCounterData, 197	counter_css_completely_false_recip_comiss, 291
$\sim$ NetCounterData, 198	counter_css_completely_false_recip_omiss, 292
indices, 198	counter_css_mixed_recip, 292
NetCounterData, 198	counter_css_partially_false_recip_commi, 292
numbers, 198	counter_css_partially_false_recip_omiss, 293
NetCounters	CSS_APPEND, 286
network.hpp, 298	CSS_CASE_ELSE, 287
NetModel	CSS_CASE_PERCEIVED, 287
network.hpp, 299	CSS_CASE_TRUTH, 287
NetRule	CSS_CHECK_SIZE, 287
network.hpp, 299	CSS_CHECK_SIZE_INIT, 287
NetRules	CSS_MATCH_TYPE, 288
network.hpp, 299	CSS_NET_COUNTER_LAMBDA_INIT, 288
NetStatsCounter	CSS_PERCEIVED_CELLS, 288
network.hpp, 299	CSS_SIZE, 288
NetSupport	CSS_TRUE_CELLS, 289
network.hpp, 299	network.hpp
Network	BARRY_ZERO_NETWORK, 296
network.hpp, 299	BARRY_ZERO_NETWORK_DENSE, 296
Network counters, 24	NET_C_DATA_IDX, 297
counter_absdiff, 26	NET_C_DATA_NUM, 297
counter_ctriads, 27	NetCounter, 298
counter_degree, 27	
	NetCounters, 298
counter_density, 27	NetCounters, 298 NetModel, 299
counter_density, 27 counter_diff, 28 counter_edges, 28	NetCounters, 298

NetStatsCounter, 299	ord, 205
NetSupport, 299	parent, 206
Network, 299	probability, 206
NETWORK_COUNTER, 297	subtree_prob, 206
NETWORK_COUNTER_LAMBDA, 297	visited, 206
NETWORK_RULE, 297	NodeData, 207
NETWORK_RULE_LAMBDA, 298	blengths, 207
NetworkDense, 300	duplication, 208
NETWORKDENSE_COUNTER_LAMBDA, 298	NodeData, 207
rules_zerodiag, 300	states, 208
NETWORK_COUNTER	nodes
Network counters, 34	Geese, 173
network.hpp, 297	noexcept
NETWORK_COUNTER_LAMBDA	counters-meat.hpp, 284
network.hpp, 297	noffspring
NETWORK_RULE	Node, 204
network.hpp, 297	NONE
NETWORK_RULE_LAMBDA	CHECK, 56
network.hpp, 298	EXISTS, 58
Network Data, 200	normalizing_constants
~NetworkData, 200	Model < Array_Type, Data_Counter_Type
directed, 200	Data_Rule_Type, Data_Rule_Dyn_Type >
NetworkData, 199, 200	193
vertex_attr, 200 NetworkDense	nrow BArray < Cell_Type, Data_Type >, 67
	BArrayDense < Cell_Type, Data_Type >, 87  BArrayDense < Cell_Type, Data_Type >, 88
network.hpp, 300 NETWORKDENSE_COUNTER_LAMBDA	Phylo rules, 41
network.hpp, 298	nrules
next	Model< Array_Type, Data_Counter_Type
Progress, 221	Data_Rule_Type, Data_Counter_Type >
nfunctions	187
Flock, 156	nrules_dyn
Geese, 173	Model< Array_Type, Data_Counter_Type
nfuns	Data_Rule_Type, Data_Rule_Dyn_Type >
Flock, 153	187
Geese, 168	nterms
nleafs	Flock, 154
Flock, 154	Geese, 168
Geese, 168	Model< Array_Type, Data_Counter_Type
nnodes	Data_Rule_Type, Data_Rule_Dyn_Type >
Flock, 154	187
Geese, 168	ntrees
nnozero	Flock, 154
BArray< Cell_Type, Data_Type >, 67	num
BArrayDense < Cell_Type, Data_Type >, 88	Phylo rules, 41, 42
Node, 201	numbers
$\sim$ Node, 203	NetCounterData, 198
annotations, 204	Phylo rules, 45, 46
array, 204	
arrays, 204	obs_start
duplication, 205	Phylo rules, 46
get_parent, 203	observed_counts
id, 205	Geese, 169
is_leaf, 204	offspring
narray, 205	Node, 205
Node, 202, 203	ONE
noffspring, 204	CHECK, 56
offspring, 205	EXISTS, 58
	operator BArrayRow $<$ Cell_Type, Data_Type $>$

BArrayRow< Cell_Type, Data_Type >, 110 operator BArrayRow_const< Cell_Type, Data_Type > BArrayRow_const< Cell_Type, Data_Type >, 112 operator Cell_Type	PhyloRuleDynData, 212 Rule < Array_Type, Data_Type >, 224 Rules < Array_Type, Data_Type >, 227 vecHasher < T >, 244
BArrayCell< Cell_Type, Data_Type >, 75 BArrayCell_const< Cell_Type, Data_Type >, 77 BArrayDenseCell< Cell_Type, Data_Type >, 97 Cell< Cell_Type >, 124	operator+= BArray< Cell_Type, Data_Type >, 68 BArrayCell< Cell_Type, Data_Type >, 75 BArrayDense< Cell_Type, Data_Type >, 89
operator std::vector< Cell_Type > BArrayVector< Cell_Type, Data_Type >, 116 BArrayVector_const< Cell_Type, Data_Type >, 119	BArrayDenseCell     Cell_Type, Data_Type >, 98       BArrayRow     Cell_Type, Data_Type >, 110       BArrayVector     Cell_Type, Data_Type >, 116       operator-=
operator!=	BArray< Cell_Type, Data_Type >, 68
BArrayCell_const< Cell_Type, Data_Type >, 77 BArrayRow_const< Cell_Type, Data_Type >, 112 BArrayVector_const< Cell_Type, Data_Type >, 119	BArrayCell< Cell_Type, Data_Type >, 75 BArrayDense< Cell_Type, Data_Type >, 90 BArrayDenseCell< Cell_Type, Data_Type >, 98 BArrayRow< Cell_Type, Data_Type >, 110
Cell< Cell_Type >, 124	BArrayVector< Cell_Type, Data_Type >, 116
operator<	operator/=
BArrayCell_const< Cell_Type, Data_Type >, 77 BArrayRow_const< Cell_Type, Data_Type >, 112 BArrayVector_const< Cell_Type, Data_Type >, 120	BArray< Cell_Type, Data_Type >, 69 BArrayCell< Cell_Type, Data_Type >, 75 BArrayDense< Cell_Type, Data_Type >, 90 BArrayDenseCell< Cell_Type, Data_Type >, 98
operator<=	BArrayRow< Cell_Type, Data_Type >, 111
BArrayCell_const< Cell_Type, Data_Type >, 78 BArrayRow_const< Cell_Type, Data_Type >, 113 BArrayVector_const< Cell_Type, Data_Type >,	BArrayVector< Cell_Type, Data_Type >, 116 operator= BArray< Cell_Type, Data_Type >, 69
120	BArrayCell< Cell_Type, Data_Type >, 75
operator>	BArrayDense< Cell_Type, Data_Type >, 90, 91
BArrayCell_const< Cell_Type, Data_Type >, 78 BArrayRow_const< Cell_Type, Data_Type >, 113 BArrayVector_const< Cell_Type, Data_Type >, 120	BArrayDenseCell     Cell_Type, Data_Type >, 98       BArrayRow     Cell_Type, Data_Type >, 111       BArrayVector     Cell_Type, Data_Type >, 117       Cell     Cell_Type >, 124, 125
operator>=	Counter< Array_Type, Data_Type >, 131, 132
BArrayCell_const< Cell_Type, Data_Type >, 78 BArrayRow_const< Cell_Type, Data_Type >, 113	Counters< Array_Type, Data_Type >, 137 Geese, 169
BArrayVector_const< Cell_Type, Data_Type >, 120	Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,
operator*= BArray< Cell_Type, Data_Type >, 67	187 Rules < Array_Type, Data_Type >, 228
BArrayCell< Cell_Type, Data_Type >, 07  BArrayCell< Cell_Type, Data_Type >, 75	operator==
BArrayDense< Cell_Type, Data_Type >, 89 BArrayDenseCell< Cell_Type, Data_Type >, 97 BArrayRow< Cell_Type, Data_Type >, 110 BArrayVector< Cell_Type, Data_Type >, 116	BArray< Cell_Type, Data_Type >, 69 BArrayCell< Cell_Type, Data_Type >, 76 BArrayCell_const< Cell_Type, Data_Type >, 78 BArrayDense< Cell_Type, Data_Type >, 91
operator()	BArrayDenseCell< Cell_Type, Data_Type >, 99
BArray< Cell_Type, Data_Type >, 67	BArrayRow< Cell_Type, Data_Type >, 111
barray-meat-operators.hpp, 249	BArrayRow_const< Cell_Type, Data_Type >, 113
BArrayDense < Cell_Type, Data_Type >, 88, 89	BArrayVector< Cell_Type, Data_Type >, 117
BArrayDenseCol< Cell_Type, Data_Type >, 101	BArrayVector_const< Cell_Type, Data_Type >,
BArrayDenseCol_const< Cell_Type, Data_Type >,	120
103	Cell< Cell_Type >, 125
BArrayDenseRow< Cell_Type, Data_Type >, 106	operator[]
BArrayDenseRow_const< Cell_Type, Data_Type	Counters < Array_Type, Data_Type >, 138
>, 108	PhyloCounterData, 210
Flock, 154	PowerSet < Array_Type, Data_Rule_Type >, 217
Phylo rules, 42 PhyloCounterData, 210	ord Node, 205
. Hylocountor Data, 210	11000, 200

out of range	DEEMBulos 39
out_of_range	DEFMRules, 38
BArray Cell_Type, Data_Type >, 69	DEFMRulesDyn, 38
BArrayDense< Cell_Type, Data_Type >, 91	DEFMStatsCounter, 38
noromo	DEFMSupport, 38
params	idx, 40, 41
model-meat.hpp, 318	indices, 44, 45
params_last	init, 45
Model Array_Type, Data_Counter_Type,	is_motif, 45
Data_Rule_Type, Data_Rule_Dyn_Type >,	is_true, 41
193	logical, 45
parent	ncol, 41
Node, 206	nrow, 41
parse_polytomies	num, 41, 42
Flock, 155	numbers, 45, 46
Geese, 169	obs_start, 46
Phylo counters, 47	operator(), 42
counter_co_opt, 48	print, 42
counter_cogain, 49	rule_dyn_limit_changes, 42
counter_gains, 49	X_ncol, 46
counter_gains_from_0, 49	X_nrow, 46
counter_gains_k_offspring, 49	PHYLO CHECK MISSING
counter_genes_changing, 50	
counter k genes changing, 50	Counting, 16
counter_less_than_p_prop_genes_changing, 50	PHYLO_COUNTER_LAMBDA
counter_longest, 50	Counting, 17
counter_loss, 51	PHYLO_RULE_DYN_LAMBDA
counter_maxfuns, 51	Counting, 17
	PhyloArray
counter_neofun, 51	geese-types.hpp, 340
counter_neofun_a2b, 51	PhyloCounter
counter_overall_changes, 52	geese-types.hpp, 340
counter_overall_gains, 52	PhyloCounterData, 208
counter_overall_gains_from_0, 52	at, 209
counter_overall_loss, 52	begin, 209
counter_pairwise_first_gain, 53	empty, 209
counter_pairwise_neofun_singlefun, 53	end, 210
counter_pairwise_overall_change, 53	get_counters, 210
counter_pairwise_preserving, 53	operator(), 210
counter_preserve_pseudogene, 54	operator[], 210
counter_prop_genes_changing, 54	PhyloCounterData, 209
counter_subfun, 54	push_back, 210
Phylo rules, 35	reserve, 210
$\sim$ DEFMCounterData, 43	shrink to fit, 211
∼DEFMData, <mark>43</mark>	size, 211
∼DEFMRuleDynData, 43	PhyloCounters
array, 44	•
at, 39	geese-types.hpp, 340
counts, 44	PhyloModel
covar_sort, 44	geese-types.hpp, 340
covar_used, 44	PhyloPowerSet
covariates, 44	geese-types.hpp, 340
DEFMCounter, 37	PhyloRule
DEFMCounterData, 39	geese-types.hpp, 341
	PhyloRuleData
DEFMCounters, 37	geese-types.hpp, 341
DEFMMadal 37	PhyloRuleDyn
DEFMModel, 37	geese-types.hpp, 341
DEFMRule, 38	PhyloRuleDynData, 211
DEFMRuleData, 40	$\sim$ PhyloRuleDynData, 212
DEFMRuleDyn, 38	counts, 212
DEFMRuleDynData, 40	

duplication 212	Gaana 170
duplication, 212	Geese, 170
lb, 213	predict_exhaust_backend
operator(), 212	Geese, 170
PhyloRuleDynData, 212	predict_sim
pos, 213	Geese, 170
ub, 213	print
PhyloRules	BArray< Cell_Type, Data_Type >, 69
geese-types.hpp, 341	BArrayDense < Cell_Type, Data_Type >, 91
PhyloRulesDyn	DEFM, 143
geese-types.hpp, 341	Flock, 155
PhyloStatsCounter	FreqTable $<$ T $>$ , 159
geese-types.hpp, 341	Geese, 170
PhyloSupport	Model< Array_Type, Data_Counter_Type,
geese-types.hpp, 342	Data_Rule_Type, Data_Rule_Dyn_Type >,
POS	188
barraydense-meat-operators.hpp, 254	Phylo rules, 42
barraydense-meat.hpp, 256	Support< Array_Type, Data_Counter_Type,
barraydensecell-bones.hpp, 258	Data_Rule_Type, Data_Rule_Dyn_Type >,
barraydensecell-meat.hpp, 258	239
barraydensecol-bones.hpp, 259	print_n
barraydenserow-bones.hpp, 261	BArray< Cell_Type, Data_Type >, 70
pos	print_nodes
PhyloRuleDynData, 213	Geese, 171
POS_N	print_observed_counts
barraydense-meat-operators.hpp, 254	Geese, 171
barraydense-meat.hpp, 257	print_stats
barraydensecol-bones.hpp, 259	Model< Array_Type, Data_Counter_Type,
barraydenserow-bones.hpp, 261	7= 71 / = = 71 /
• • • • • • • • • • • • • • • • • • • •	Data_Rule_Type, Data_Rule_Dyn_Type >,
PowerSet	188
PowerSet< Array_Type, Data_Rule_Type >, 215	printf_barry
PowerSet < Array_Type, Data_Rule_Type >, 213	barry-configuration.hpp, 265
$\sim$ PowerSet, 215	probability
add_rule, 215, 216	Node, 206
begin, 216	probs
calc, 216	model-meat.hpp, 318
coordinates_free, 218	
	Progress, 220
coordinates locked, 218	
coordinates_locked, 218 data, 218	∼Progress, 220
data, 218	∼Progress, 220 end, 221
data, 218 EmptyArray, 218	~Progress, 220 end, 221 next, 221
data, 218 EmptyArray, 218 end, 216	~Progress, 220 end, 221 next, 221 Progress, 220
data, 218 EmptyArray, 218 end, 216 get_data, 216	~Progress, 220 end, 221 next, 221 Progress, 220 progress.hpp
data, 218 EmptyArray, 218 end, 216 get_data, 216 get_data_ptr, 217	~Progress, 220 end, 221 next, 221 Progress, 220 progress.hpp BARRY_PROGRESS_BAR_WIDTH, 344
data, 218 EmptyArray, 218 end, 216 get_data, 216 get_data_ptr, 217 init_support, 217	~Progress, 220 end, 221 next, 221 Progress, 220 progress.hpp BARRY_PROGRESS_BAR_WIDTH, 344 pset_arrays
data, 218 EmptyArray, 218 end, 216 get_data, 216 get_data_ptr, 217 init_support, 217 M, 218	~Progress, 220 end, 221 next, 221 Progress, 220 progress.hpp BARRY_PROGRESS_BAR_WIDTH, 344 pset_arrays Model< Array_Type, Data_Counter_Type,
data, 218 EmptyArray, 218 end, 216 get_data, 216 get_data_ptr, 217 init_support, 217 M, 218 N, 219	~Progress, 220 end, 221 next, 221 Progress, 220 progress.hpp BARRY_PROGRESS_BAR_WIDTH, 344 pset_arrays Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,
data, 218 EmptyArray, 218 end, 216 get_data, 216 get_data_ptr, 217 init_support, 217 M, 218	~Progress, 220 end, 221 next, 221 Progress, 220 progress.hpp BARRY_PROGRESS_BAR_WIDTH, 344 pset_arrays Model< Array_Type, Data_Counter_Type,
data, 218 EmptyArray, 218 end, 216 get_data, 216 get_data_ptr, 217 init_support, 217 M, 218 N, 219	~Progress, 220 end, 221 next, 221 Progress, 220 progress.hpp BARRY_PROGRESS_BAR_WIDTH, 344 pset_arrays Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,
data, 218 EmptyArray, 218 end, 216 get_data, 216 get_data_ptr, 217 init_support, 217 M, 218 N, 219 n_free, 219	~Progress, 220 end, 221 next, 221 Progress, 220 progress.hpp BARRY_PROGRESS_BAR_WIDTH, 344 pset_arrays Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 193
data, 218 EmptyArray, 218 end, 216 get_data, 216 get_data_ptr, 217 init_support, 217 M, 218 N, 219 n_free, 219 n_locked, 219	~Progress, 220 end, 221 next, 221 Progress, 220 progress.hpp BARRY_PROGRESS_BAR_WIDTH, 344 pset_arrays Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 193 model-meat.hpp, 318
data, 218 EmptyArray, 218 end, 216 get_data, 216 get_data_ptr, 217 init_support, 217 M, 218 N, 219 n_free, 219 n_locked, 219 operator[], 217 PowerSet, 215	~Progress, 220 end, 221 next, 221 Progress, 220 progress.hpp BARRY_PROGRESS_BAR_WIDTH, 344 pset_arrays Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 193 model-meat.hpp, 318 pset_loc Geese, 173
data, 218 EmptyArray, 218 end, 216 get_data, 216 get_data_ptr, 217 init_support, 217 M, 218 N, 219 n_free, 219 n_locked, 219 operator[], 217 PowerSet, 215 reset, 217	~Progress, 220 end, 221 next, 221 Progress, 220 progress.hpp BARRY_PROGRESS_BAR_WIDTH, 344 pset_arrays Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 193 model-meat.hpp, 318 pset_loc Geese, 173 pset_probs
data, 218 EmptyArray, 218 end, 216 get_data, 216 get_data_ptr, 217 init_support, 217 M, 218 N, 219 n_free, 219 n_locked, 219 operator[], 217 PowerSet, 215 reset, 217 rules, 219	~Progress, 220 end, 221 next, 221 Progress, 220 progress.hpp BARRY_PROGRESS_BAR_WIDTH, 344 pset_arrays Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 193 model-meat.hpp, 318 pset_loc Geese, 173 pset_probs Model< Array_Type, Data_Counter_Type,
data, 218 EmptyArray, 218 end, 216 get_data, 216 get_data_ptr, 217 init_support, 217 M, 218 N, 219 n_free, 219 n_locked, 219 operator[], 217 PowerSet, 215 reset, 217 rules, 219 rules_deleted, 219	~Progress, 220 end, 221 next, 221 Progress, 220 progress.hpp BARRY_PROGRESS_BAR_WIDTH, 344 pset_arrays Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 193 model-meat.hpp, 318 pset_loc Geese, 173 pset_probs Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,
data, 218 EmptyArray, 218 end, 216 get_data, 216 get_data_ptr, 217 init_support, 217 M, 218 N, 219 n_free, 219 n_locked, 219 operator[], 217 PowerSet, 215 reset, 217 rules, 219 rules_deleted, 219 size, 217	~Progress, 220 end, 221 next, 221 Progress, 220 progress.hpp BARRY_PROGRESS_BAR_WIDTH, 344 pset_arrays Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 193 model-meat.hpp, 318 pset_loc Geese, 173 pset_probs Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 194
data, 218 EmptyArray, 218 end, 216 get_data, 216 get_data_ptr, 217 init_support, 217 M, 218 N, 219 n_free, 219 n_locked, 219 operator[], 217 PowerSet, 215 reset, 217 rules, 219 rules_deleted, 219 size, 217 predict	~Progress, 220 end, 221 next, 221 Progress, 220 progress.hpp BARRY_PROGRESS_BAR_WIDTH, 344 pset_arrays Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 193 model-meat.hpp, 318 pset_loc Geese, 173 pset_probs Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 194 pset_stats
data, 218 EmptyArray, 218 end, 216 get_data, 216 get_data_ptr, 217 init_support, 217 M, 218 N, 219 n_free, 219 n_locked, 219 operator[], 217 PowerSet, 215 reset, 217 rules, 219 rules_deleted, 219 size, 217 predict Geese, 169	~Progress, 220 end, 221 next, 221 Progress, 220 progress.hpp BARRY_PROGRESS_BAR_WIDTH, 344 pset_arrays Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 193 model-meat.hpp, 318 pset_loc Geese, 173 pset_probs Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 194 pset_stats Model< Array_Type, Data_Counter_Type, Data_Stats Model< Array_Type, Data_Counter_Type,
data, 218 EmptyArray, 218 end, 216 get_data, 216 get_data_ptr, 217 init_support, 217 M, 218 N, 219 n_free, 219 n_locked, 219 operator[], 217 PowerSet, 215 reset, 217 rules, 219 rules_deleted, 219 size, 217 predict Geese, 169 predict_backend	~Progress, 220 end, 221 next, 221 Progress, 220 progress.hpp BARRY_PROGRESS_BAR_WIDTH, 344 pset_arrays Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 193 model-meat.hpp, 318 pset_loc Geese, 173 pset_probs Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 194 pset_stats Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 194 pset_stats Model< Array_Type, Data_Rule_Dyn_Type >, Data_Rule_Type, Data_Rule_Dyn_Type >,
data, 218 EmptyArray, 218 end, 216 get_data, 216 get_data_ptr, 217 init_support, 217 M, 218 N, 219 n_free, 219 n_locked, 219 operator[], 217 PowerSet, 215 reset, 217 rules, 219 rules_deleted, 219 size, 217 predict Geese, 169	~Progress, 220 end, 221 next, 221 Progress, 220 progress.hpp BARRY_PROGRESS_BAR_WIDTH, 344 pset_arrays Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 193 model-meat.hpp, 318 pset_loc Geese, 173 pset_probs Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 194 pset_stats Model< Array_Type, Data_Counter_Type, Data_Stats Model< Array_Type, Data_Counter_Type,

model-meat.hpp, 312	get_description, 223
PhyloCounterData, 210	get_name, 223
r	operator(), 224
model-meat.hpp, 318	Rule, 222
README.md, 358	rule_dyn_limit_changes
reduced_sequence	Phylo rules, 42
Geese, 174	rule_fun_
rengine	model-meat.hpp, 318 rule_fun_default
Flock, 156	rules-bones.hpp, 344
Model < Array_Type, Data_Counter_Type,	Rule fun type
Data_Rule_Type, Data_Rule_Dyn_Type >,	typedefs.hpp, 356
194	RULE_FUNCTION
res	barry.hpp, 271
counters-meat.hpp, 284	geese-bones.hpp, 333
reserve	RULE LAMBDA
BArray< Cell_Type, Data_Type >, 70	barry.hpp, 271
BArrayDense< Cell_Type, Data_Type >, 91	Rules
FreqTable $<$ T $>$ , 159	Rules< Array_Type, Data_Type >, 225
model-meat.hpp, 312	rules
PhyloCounterData, 210	Model< Array_Type, Data_Counter_Type,
reset	Data_Rule_Type, Data_Rule_Dyn_Type >,
PowerSet < Array_Type, Data_Rule_Type >, 217	194
reset_array	model-meat.hpp, 319
StatsCounter< Array_Type, Data_Type >, 231	PowerSet < Array_Type, Data_Rule_Type >, 219
Support< Array_Type, Data_Counter_Type,	Rules< Array_Type, Data_Type >, 224
Data_Rule_Type, Data_Rule_Dyn_Type >,	$\sim$ Rules, 225
239	add_rule, 226
resize	begin, 226
BArray Cell_Type, Data_Type >, 70	end, 226
BArrayDense < Cell_Type, Data_Type >, 92	get_descriptions, 226
Entries < Cell_Type >, 149	get_names, 227
statscounter-meat.hpp, 348	get_seq, 227
return	operator(), 227
counters-meat.hpp, 285	operator=, 228
model-meat.hpp, 313, 318 statscounter-meat.hpp, 351	Rules, 225
rhs	size, 228
barray-meat-operators.hpp, 249	rules-bones.hpp
rm_cell	rule_fun_default, 344
BArray< Cell Type, Data Type >, 70	rules_
BArrayDense< Cell_Type, Data_Type >, 92	model-meat.hpp, 319
ROW	rules_deleted PowerSet< Array Type, Data Rule Type >, 219
barray-meat-operators.hpp, 247	rules dont become zero
barray-meat.hpp, 251	Network counters, 34
barraydense-meat-operators.hpp, 254	rules_dyn
barraydense-meat.hpp, 257	Model< Array_Type, Data_Counter_Type,
row	Data_Rule_Type, Data_Rule_Dyn_Type >,
BArray< Cell_Type, Data_Type >, 70	195
BArrayDense < Cell_Type, Data_Type >, 92	model-meat.hpp, 319
Row_type	rules_markov_fixed
typedefs.hpp, 356	Network counters, 35
rowsum	rules_zerodiag
BArrayDense < Cell_Type, Data_Type >, 92	network.hpp, 300
Rule	117
Rule < Array_Type, Data_Type >, 222	sample
Rule < Array_Type, Data_Type >, 221	Model < Array_Type, Data_Counter_Type,
$\sim$ Rule, 222	$Data_Rule_Type, Data_Rule_Dyn_Type >,$
D, 223	188

sequence	BArrayDenseRow_const< Cell_Type, Data_Type
Geese, 174	>, 108
set_counters	BArrayVector< Cell_Type, Data_Type >, 117
Model< Array_Type, Data_Counter_Type,	BArrayVector_const< Cell_Type, Data_Type >,
Data_Rule_Type, Data_Rule_Dyn_Type >,	121
189	Counters< Array_Type, Data_Type >, 138
model-meat.hpp, 313	FreqTable $< T >$ , 160
StatsCounter< Array_Type, Data_Type >, 232	Model< Array_Type, Data_Counter_Type,
Support< Array_Type, Data_Counter_Type,	Data_Rule_Type, Data_Rule_Dyn_Type >,
Data_Rule_Type, Data_Rule_Dyn_Type >,	190
239	model-meat.hpp, 313
	PhyloCounterData, 211
Set_data  PArroy < Coll Type Data Type > 71	-
BArray Penas ( Cell Time Pata Time > 00	PowerSet < Array_Type, Data_Rule_Type >, 217
BArrayDense < Cell_Type, Data_Type >, 93	Rules < Array_Type, Data_Type >, 228
set_hasher	StatsCounter< Array_Type, Data_Type >, 232
Counter< Array_Type, Data_Type >, 132	size_unique
set_names	Model< Array_Type, Data_Counter_Type,
DEFM, 143	Data_Rule_Type, Data_Rule_Dyn_Type >,
set_rengine	190
Model< Array_Type, Data_Counter_Type,	sort_array
Data_Rule_Type, Data_Rule_Dyn_Type >,	typedefs.hpp, 356
189	source
set_rules	Entries < Cell_Type >, 149
Model< Array_Type, Data_Counter_Type,	states
Data_Rule_Type, Data_Rule_Dyn_Type >,	NodeData, 208
189	Statistical Models, 24
model-meat.hpp, 313	stats
Support< Array_Type, Data_Counter_Type,	model-meat.hpp, 319
Data_Rule_Type, Data_Rule_Dyn_Type >,	stats_support
240	Model< Array_Type, Data_Counter_Type,
set_rules_dyn	Data_Rule_Type, Data_Rule_Dyn_Type >,
Model< Array_Type, Data_Counter_Type,	195
Data_Rule_Type, Data_Rule_Dyn_Type >,	stats support n arrays
189	Model Array_Type, Data_Counter_Type,
model-meat.hpp, 313	Data_Rule_Type, Data_Rule_Dyn_Type >,
Support< Array_Type, Data_Counter_Type,	195
Data_Rule_Type, Data_Rule_Dyn_Type >,	model-meat.hpp, 319
240	stats target
set seed	Model< Array_Type, Data_Counter_Type,
Flock, 155	Data_Rule_Type, Data_Rule_Dyn_Type >,
Geese, 171	195
Model< Array_Type, Data_Counter_Type,	StatsCounter
Data_Rule_Type, Data_Rule_Dyn_Type >,	StatsCounter< Array_Type, Data_Type >, 229,
189	230
set transform model	StatsCounter< Array_Type, Data_Type >, 228
Model< Array_Type, Data_Counter_Type,	~StatsCounter, 230
Data_Rule_Type, Data_Rule_Dyn_Type >,	add counter, 230
190	count all, 230
shrink_to_fit	count_current, 231
PhyloCounterData, 211	count_init, 231
simulate	get_counters, 231
DEFM, 143	get_descriptions, 231
Geese, 171	get_names, 231
Size  RArrayDansaCol / Call Type Data Type > 101	reset_array, 231 set_counters, 232
BArrayDenseCol< Cell_Type, Data_Type >, 101 BArrayDenseCol_const< Cell_Type, Data_Type >,	
104	size, 232 StateCounter, 229, 230
	StatsCounter, 229, 230
BArrayDenseRow< Cell_Type, Data_Type >, 106	statscounter-meat.hpp

clear, 347	max_num_elements, 243
counter, 349	N, 243
counter_deleted, 350	n_counters, 243
counters, 350	print, 239
counters_, 350	reset_array, 239
current_stats, 350	set_counters, 239
EmptyArray, 350	set_rules, 240
f_, 351	set_rules_dyn, 240
for, 347	Support, 234, 235
j, 351	support-meat.hpp
resize, 348 return, 351	BARRY_SUPPORT_MEAT_HPP, 352 support_fun
STATSCOUNTER_TEMPLATE, 347–349	Model Array_Type, Data_Counter_Type,
STATSCOUNTER_TEMPLATE_ARGS, 347	Data_Rule_Type, Data_Rule_Dyn_Type >,
STATSCOUNTER_TYPE, 347	196
STATSCOUNTER_TEMPLATE	support_size
statscounter-meat.hpp, 347–349	Flock, 155
STATSCOUNTER_TEMPLATE_ARGS	Geese, 171
statscounter-meat.hpp, 347	Model < Array_Type, Data_Counter_Type,
STATSCOUNTER_TYPE	Data_Rule_Type, Data_Rule_Dyn_Type >,
statscounter-meat.hpp, 347	191
store_psets	swap_cells
Model< Array_Type, Data_Counter_Type,	BArray< Cell_Type, Data_Type >, 71
Data_Rule_Type, Data_Rule_Dyn_Type >,	BArrayDense< Cell_Type, Data_Type >, 93
190	swap_cols
subtree_prob	BArray< Cell_Type, Data_Type >, 71
Node, 206	BArrayDense < Cell_Type, Data_Type >, 93
Support Support Time Date Country Time	swap_rows
Support	BArray Cell_Type, Data_Type >, 71
Data_Rule_Type, Data_Rule_Dyn_Type >, 234, 235	BArrayDense< Cell_Type, Data_Type >, 93
Support< Array_Type, Data_Counter_Type, Data_Rule_T	-, target
Data_Rule_Dyn_Type >, 232	Entries < Cell_Type >, 149
~Support, 235	temp_stats
add_counter, 235	model-meat.hpp, 320
add_rule, 235, 236	this
add_rule_dyn, 236	barray-meat-operators.hpp, 250
calc, 236	tmp_counts
change_stats, 240	model-meat.hpp, 320
coordiantes_n_free, 240	TMP_HASHER_CALL
coordiantes_n_locked, 241	counters-meat.hpp, 275
coordinates_free, 241	toggle_cell
coordinates_locked, 241	BArray Cell_Type, Data_Type >, 72
current_stats, 241	BArrayDense < Cell_Type, Data_Type >, 94 toggle_lock
delete_counters, 241	BArray< Cell Type, Data Type >, 72
delete_rules, 242	BArrayDense< Cell_Type, Data_Type >, 72
delete_rules_dyn, 242	transform_model
eval_rules_dyn, 237	Model< Array_Type, Data_Counter_Type,
get_counters, 237 get_counts, 237	Data_Rule_Type, Data_Rule_Dyn_Type >,
get_current_stats, 237	191
get_data, 238	transform_model_fun
get_rules, 238	Model< Array_Type, Data_Counter_Type,
get_rules_dyn, 238	Data_Rule_Type, Data_Rule_Dyn_Type >,
hashes, 242	196
hashes_initialized, 242	transform_model_term_names
init_support, 238	Model< Array_Type, Data_Counter_Type,
M, 242	Data_Rule_Type, Data_Rule_Dyn_Type >, 196

```
Model<
                                                                                      Data_Counter_Type,
transpose
                                                                      Array_Type,
     BArray< Cell_Type, Data_Type >, 72
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
    BArrayDense < Cell_Type, Data_Type >, 94
                                                               197
TWO
                                                     X ncol
    CHECK, 56
                                                          Phylo rules, 46
    EXISTS, 58
                                                     X nrow
typedefs.hpp
                                                          Phylo rules, 46
    Col type, 355
    Counter_fun_type, 355
                                                     ZERO CELL
    Counts_type, 355
                                                          barraydense-meat.hpp, 257
    Hasher_fun_type, 355
                                                          barraydensecol-bones.hpp, 260
    MapVec_type, 356
                                                          barraydenserow-bones.hpp, 261
    Row_type, 356
                                                     zero col
    Rule_fun_type, 356
                                                          BArray< Cell_Type, Data_Type >, 72
    sort_array, 356
                                                          BArrayDense < Cell_Type, Data_Type >, 94
    vec_equal, 357
                                                     zero row
    vec_equal_approx, 357
                                                          BArray< Cell_Type, Data_Type >, 72
    vec inner prod, 357, 358
                                                          BArrayDense < Cell_Type, Data_Type >, 94
ub
     PhyloRuleDynData, 213
UKNOWN
    EXISTS, 58
UNI SUB
    counters.hpp, 323
update annotations
    Geese, 171
update_normalizing_constant
    model-meat.hpp, 313
urand
    model-meat.hpp, 314
val
     Entries < Cell_Type >, 149
value
    Cell< Cell_Type >, 125
vec diff
    geese-bones.hpp, 333
vec_equal
    typedefs.hpp, 357
vec_equal_approx
    typedefs.hpp, 357
vec inner prod
    typedefs.hpp, 357, 358
vecHasher< T >, 243
    operator(), 244
vector caster
    geese-bones.hpp, 334
vertex_attr
    NetworkData, 200
visited
    BArray< Cell_Type, Data_Type >, 73
    BArrayDense < Cell Type, Data Type >, 96
    Cell< Cell_Type >, 125
    Node, 206
with pset
```