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
8.29 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class	
plate Reference	
8.29.1 Detailed Description	
8.29.2 Constructor & Destructor Documentation	
8.29.2.1 Model() [1/3]	
<b>8.29.2.2 Model()</b> [2/3]	
<b>8.29.2.3 Model()</b> [3/3]	
8.29.2.4 ~Model()	
8.29.3 Member Function Documentation	
8.29.3.1 add_array()	 . 180
8.29.3.2 add_counter() [1/2]	 . 181
8.29.3.3 add_counter() [2/2]	
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()	 . 184
8.29.3.15 get_pset()	 . 184
8.29.3.16 get_pset_arrays()	 . 184
8.29.3.17 get_pset_probs()	 . 184
<b>8.29.3.18</b> get_pset_stats() [1/2]	 . 185
<b>8.29.3.19</b> get_pset_stats() [2/2]	 . 185
8.29.3.20 get_rengine()	 . 185
8 29 3 21 get_rules()	185

8.29.3.22 get_rules_dyn()	186
8.29.3.23 get_stats_support()	
8.29.3.24 get_stats_target()	
8.29.3.25 get_support_fun()	
8.29.3.26 likelihood() [1/4]	
8.29.3.27 likelihood() [2/4]	187
8.29.3.28 likelihood() [3/4]	
8.29.3.29 likelihood() [4/4]	187
8.29.3.30 likelihood_total()	188
8.29.3.31 nrules()	188
8.29.3.32 nrules_dyn()	188
8.29.3.33 nterms()	188
8.29.3.34 operator=()	189
8.29.3.35 print()	189
8.29.3.36 print_stats()	189
8.29.3.37 sample() [1/2]	189
8.29.3.38 sample() [2/2]	190
8.29.3.39 set_counters()	190
8.29.3.40 set_rengine()	190
8.29.3.41 set_rules()	190
8.29.3.42 set_rules_dyn()	191
8.29.3.43 set_seed()	191
8.29.3.44 set_transform_model()	191
8.29.3.45 size()	191
8.29.3.46 size_unique()	192
8.29.3.47 store_psets()	192
8.29.3.48 support_size()	192
8.29.3.49 transform_model()	192
8.29.4 Member Data Documentation	193
8.29.4.1 arrays2support	193
8.29.4.2 counter_fun	193
8.29.4.3 counters	193
8.29.4.4 delete_counters	193
8.29.4.5 delete_rengine	194
8.29.4.6 delete_rules	194
8.29.4.7 delete_rules_dyn	194
8.29.4.8 first_calc_done	194
8.29.4.9 keys2support	194
8.29.4.10 normalizing_constants	
8.29.4.11 params_last	
8.29.4.12 pset_arrays	195
8.29.4.13 pset_probs	195

8.29.4.14 pset_stats	196
8.29.4.15 rengine	196
8.29.4.16 rules	196
8.29.4.17 rules_dyn	196
8.29.4.18 stats_support	197
8.29.4.19 stats_support_n_arrays	197
8.29.4.20 stats_target	197
8.29.4.21 support_fun	197
8.29.4.22 transform_model_fun	198
8.29.4.23 transform_model_term_names	198
8.29.4.24 with_pset	198
8.30 NetCounterData Class Reference	199
8.30.1 Detailed Description	199
8.30.2 Constructor & Destructor Documentation	199
8.30.2.1 NetCounterData() [1/2]	199
8.30.2.2 NetCounterData() [2/2]	199
8.30.2.3 ~NetCounterData()	200
8.30.3 Member Data Documentation	200
8.30.3.1 indices	200
8.30.3.2 numbers	200
8.31 NetworkData Class Reference	200
8.31.1 Detailed Description	201
8.31.2 Constructor & Destructor Documentation	201
8.31.2.1 NetworkData() [1/3]	201
8.31.2.2 NetworkData() [2/3]	201
<b>8.31.2.3 NetworkData()</b> [3/3]	201
8.31.2.4 ~NetworkData()	202
8.31.3 Member Data Documentation	202
8.31.3.1 directed	202
8.31.3.2 vertex_attr	202
8.32 Node Class Reference	202
8.32.1 Detailed Description	203
8.32.2 Constructor & Destructor Documentation	203
<b>8.32.2.1 Node()</b> [1/5]	204
<b>8.32.2.2 Node()</b> [2/5]	204
<b>8.32.2.3 Node()</b> [3/5]	204
<b>8.32.2.4 Node()</b> [4/5]	204
<b>8.32.2.5 Node()</b> [5/5]	204
8.32.2.6 ~Node()	205
8.32.3 Member Function Documentation	205
8.32.3.1 get_parent()	205
8 32 3 2 is leaf()	205

8.32.3.3 noffspring()	05
8.32.4 Member Data Documentation	05
8.32.4.1 annotations	05
8.32.4.2 array	06
8.32.4.3 arrays	06
8.32.4.4 duplication	06
8.32.4.5 id	06
8.32.4.6 narray	06
8.32.4.7 offspring	07
8.32.4.8 ord	07
8.32.4.9 parent	07
8.32.4.10 probability	07
8.32.4.11 subtree_prob	07
8.32.4.12 visited	80
8.33 NodeData Class Reference	80
8.33.1 Detailed Description	80
8.33.2 Constructor & Destructor Documentation	80
8.33.2.1 NodeData()	80
8.33.3 Member Data Documentation	09
8.33.3.1 blengths	09
8.33.3.2 duplication	09
8.33.3.3 states	09
8.34 PhyloCounterData Class Reference	09
8.34.1 Detailed Description	10
8.34.2 Constructor & Destructor Documentation	10
8.34.2.1 PhyloCounterData() [1/2]	10
8.34.2.2 PhyloCounterData() [2/2]	10
8.34.3 Member Function Documentation	10
8.34.3.1 at()	10
8.34.3.2 begin()	10
8.34.3.3 empty()	11
8.34.3.4 end()	11
8.34.3.5 get_counters()	11
8.34.3.6 operator()()	11
8.34.3.7 operator[]()	11
8.34.3.8 push_back()	11
8.34.3.9 reserve()	12
8.34.3.10 shrink_to_fit()	12
8.34.3.11 size()	12
8.35 PhyloRuleDynData Class Reference	12
8.35.1 Detailed Description	13
8 35 2 Constructor & Destructor Documentation	12

8.35.2.1 PhyloRuleDynData()	13
8.35.2.2 $\sim$ PhyloRuleDynData()	13
8.35.3 Member Function Documentation	13
8.35.3.1 operator()()	13
8.35.4 Member Data Documentation	13
8.35.4.1 counts	13
8.35.4.2 duplication	14
8.35.4.3 lb	214
8.35.4.4 pos	14
8.35.4.5 ub	214
8.36 PowerSet< Array_Type, Data_Rule_Type > Class Template Reference	<u>'</u> 14
8.36.1 Detailed Description	15
8.36.2 Constructor & Destructor Documentation	16
8.36.2.1 PowerSet() [1/3]	16
8.36.2.2 PowerSet() [2/3]	216
8.36.2.3 PowerSet() [3/3]	16
8.36.2.4 ~PowerSet()	216
8.36.3 Member Function Documentation	16
8.36.3.1 add_rule() [1/2]	17
8.36.3.2 add_rule() [2/2]	17
8.36.3.3 begin()	17
8.36.3.4 calc()	217
8.36.3.5 end()	217
8.36.3.6 get_data()	18
8.36.3.7 get_data_ptr()	18
8.36.3.8 init_support()	18
8.36.3.9 operator[]()	18
8.36.3.10 reset()	18
8.36.3.11 size()	19
8.36.4 Member Data Documentation	19
8.36.4.1 coordinates_free	19
8.36.4.2 coordinates_locked	19
8.36.4.3 data	19
8.36.4.4 EmptyArray	19
8.36.4.5 M	20
8.36.4.6 N	20
8.36.4.7 n_free	20
8.36.4.8 n_locked	20
8.36.4.9 rules	20
8.36.4.10 rules_deleted	21
8.37 Progress Class Reference	21
8 37 1 Detailed Description	21

8.37.2 Constructor & Destructor Documentation	221
8.37.2.1 Progress()	221
8.37.2.2 ~Progress()	222
8.37.3 Member Function Documentation	222
8.37.3.1 end()	222
8.37.3.2 next()	222
8.38 Rule < Array_Type, Data_Type > Class Template Reference	
8.38.1 Detailed Description	
8.38.2 Constructor & Destructor Documentation	
8.38.2.1 Rule() [1/2]	
8.38.2.2 Rule() [2/2]	223
8.38.2.3 ~Rule()	224
8.38.3 Member Function Documentation	224
8.38.3.1 D()	224
8.38.3.2 get_description() [1/2]	224
8.38.3.3 get_description() [2/2]	224
<b>8.38.3.4 get_name()</b> [1/2]	224
8.38.3.5 get_name() [2/2]	225
8.38.3.6 operator()()	225
8.39 Rules < Array_Type, Data_Type > Class Template Reference	225
8.39.1 Detailed Description	226
8.39.2 Constructor & Destructor Documentation	226
8.39.2.1 Rules() [1/2]	226
8.39.2.2 Rules() [2/2]	226
8.39.2.3 ~Rules()	227
8.39.3 Member Function Documentation	227
8.39.3.1 add_rule() [1/2]	227
<b>8.39.3.2 add_rule()</b> [2/2]	227
8.39.3.3 begin()	227
8.39.3.4 end()	227
8.39.3.5 get_descriptions()	228
8.39.3.6 get_names()	228
8.39.3.7 get_seq()	228
8.39.3.8 operator()()	228
8.39.3.9 operator=()	229
8.39.3.10 size()	229
8.40 StatsCounter< Array_Type, Data_Type > Class Template Reference	229
8.40.1 Detailed Description	230
8.40.2 Constructor & Destructor Documentation	230
8.40.2.1 StatsCounter() [1/3]	230
8.40.2.2 StatsCounter() [2/3]	231
8.40.2.3 StatsCounter() [3/3]	

8.40.2.4 ~StatsCounter()	 231
8.40.3 Member Function Documentation	 231
8.40.3.1 add_counter()	 231
8.40.3.2 count_all()	 232
8.40.3.3 count_current()	 232
8.40.3.4 count_init()	 232
8.40.3.5 get_counters()	 232
8.40.3.6 get_descriptions()	 232
8.40.3.7 get_names()	 232
8.40.3.8 reset_array()	 232
8.40.3.9 set_counters()	 233
8.40.3.10 size()	 233
8.41 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class	
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]	 236
<b>8.41.2.3 Support()</b> [3/3]	 236
8.41.2.4 ~Support()	 236
8.41.3 Member Function Documentation	 236
8.41.3.1 add_counter()	 236
<b>8.41.3.2</b> add_rule() [1/2]	 237
<b>8.41.3.3 add_rule()</b> [2/2]	 237
<b>8.41.3.4 add_rule_dyn()</b> [1/2]	 237
<b>8.41.3.5</b> add_rule_dyn() [2/2]	 237
8.41.3.6 calc()	 237
8.41.3.7 eval_rules_dyn()	 238
8.41.3.8 get_counters()	 238
8.41.3.9 get_counts()	 238
8.41.3.10 get_current_stats()	 239
8.41.3.11 get_data()	 239
8.41.3.12 get_rules()	 239
8.41.3.13 get_rules_dyn()	 239
8.41.3.14 init_support()	 240
8.41.3.15 print()	 240
8.41.3.16 reset_array() [1/2]	 240
8.41.3.17 reset_array() [2/2]	 240
8.41.3.18 set_counters()	 241
8.41.3.19 set_rules()	 241
8.41.3.20 set_rules_dyn()	 241
8.41.4 Member Data Documentation	2/1

8.41.4.1 change_stats	 241
8.41.4.2 coordiantes_n_free	 242
8.41.4.3 coordiantes_n_locked	 242
8.41.4.4 coordinates_free	 242
8.41.4.5 coordinates_locked	 242
8.41.4.6 current_stats	 242
8.41.4.7 delete_counters	 243
8.41.4.8 delete_rules	 243
8.41.4.9 delete_rules_dyn	 243
8.41.4.10 hashes	 243
8.41.4.11 hashes_initialized	 243
8.41.4.12 M	 244
8.41.4.13 max_num_elements	 244
8.41.4.14 N	 244
8.41.4.15 n_counters	 244
8.42 vecHasher < T > Struct Template Reference	 244
8.42.1 Detailed Description	 245
8.42.2 Member Function Documentation	 245
8.42.2.1 operator()()	 245
9 File Documentation	247
9.1 include/barry/barray-bones.hpp File Reference	
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	 249
9.3.2 Function Documentation	 249
9.3.2.1 BARRAY_TEMPLATE() [1/6]	 250
9.3.2.2 BARRAY_TEMPLATE() [2/6]	
9.3.2.3 BARRAY_TEMPLATE() [3/6]	 250
9.3.2.4 BARRAY_TEMPLATE() [4/6]	 250
9.3.2.5 BARRAY_TEMPLATE() [5/6]	 250
9.3.2.6 BARRAY_TEMPLATE() [6/6]	 251
9.3.2.7 BARRAY_TEMPLATE_ARGS()	 251
9.3.2.7 BARRAY_TEMPLATE_ARGS()	
	 251
9.3.2.8 BARRAY_TYPE()	 251 251
9.3.2.8 BARRAY_TYPE()	 251 251 251

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 26

9.12.1.3 ZERO_CELL
9.13 include/barry/barraydenserow-bones.hpp File Reference
9.13.1 Macro Definition Documentation
9.13.1.1 POS
9.13.1.2 POS_N
9.13.1.3 ZERO_CELL
9.14 include/barry/barrayrow-bones.hpp File Reference
9.15 include/barry/barrayrow-meat.hpp File Reference
9.15.1 Macro Definition Documentation
9.15.1.1 BROW_TEMPLATE
9.15.1.2 BROW_TEMPLATE_ARGS
9.15.1.3 BROW_TYPE
9.15.2 Function Documentation
9.15.2.1 BROW_TEMPLATE() [1/5]
9.15.2.2 BROW_TEMPLATE() [2/5]
9.15.2.3 BROW_TEMPLATE() [3/5]
9.15.2.4 BROW_TEMPLATE() [4/5]
9.15.2.5 BROW_TEMPLATE() [5/5]
9.16 include/barry/barrayvector-bones.hpp File Reference
9.17 include/barry/barrayvector-meat.hpp File Reference
9.18 include/barry/barry-configuration.hpp File Reference
9.18.1 Macro Definition Documentation
9.18.1.1 BARRY_CHECK_SUPPORT
9.18.1.2 BARRY_ISFINITE
9.18.1.3 BARRY_MAX_NUM_ELEMENTS
9.18.1.4 BARRY_SAFE_EXP
9.18.1.5 printf_barry
9.18.2 Typedef Documentation
9.18.2.1 Map
9.19 include/barry/barry-debug.hpp File Reference
9.19.1 Macro Definition Documentation
9.19.1.1 BARRY_DEBUG_LEVEL
9.20 include/barry/barry-macros.hpp File Reference
9.20.1 Macro Definition Documentation
9.20.1.1 BARRY_NCORES_ARG
9.20.1.2 BARRY_ONE
9.20.1.3 BARRY_ONE_DENSE
9.20.1.4 BARRY_UNUSED
9.20.1.5 BARRY_ZERO
9.20.1.6 BARRY_ZERO_DENSE
9.21 include/barry/barry.hpp File Reference
9.21.1 Macro Definition Documentation

9.21.1.1 BARRY_HPP
9.21.1.2 BARRY_VERSION
9.21.1.3 BARRY_VERSION_MAYOR
9.21.1.4 BARRY_VERSION_MINOR
9.21.1.5 COUNTER_FUNCTION
9.21.1.6 COUNTER_LAMBDA
9.21.1.7 RULE_FUNCTION
9.21.1.8 RULE_LAMBDA
9.22 include/barry/cell-bones.hpp File Reference
9.23 include/barry/cell-meat.hpp File Reference
9.24 include/barry/col-bones.hpp File Reference
9.25 include/barry/counters-bones.hpp File Reference
9.26 include/barry/counters-meat.hpp File Reference
9.26.1 Macro Definition Documentation
9.26.1.1 COUNTER_TEMPLATE
9.26.1.2 COUNTER_TEMPLATE_ARGS
9.26.1.3 COUNTER_TYPE
9.26.1.4 COUNTERS_TEMPLATE
9.26.1.5 COUNTERS_TEMPLATE_ARGS
9.26.1.6 COUNTERS_TYPE
9.26.1.7 TMP_HASHER_CALL
9.26.2 Function Documentation
9.26.2.1 count_fun()
9.26.2.2 COUNTER_TEMPLATE() [1/9]
9.26.2.3 COUNTER_TEMPLATE() [2/9]
<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]
<b>9.26.2.13 COUNTERS_TEMPLATE()</b> [3/9]
9.26.2.14 COUNTERS_TEMPLATE() [4/9]
<b>9.26.2.15 COUNTERS_TEMPLATE()</b> [5/9]
9.26.2.16 COUNTERS_TEMPLATE() [6/9]
<b>9.26.2.17 COUNTERS_TEMPLATE()</b> [7/9]
9.26.2.18 COUNTERS_TEMPLATE() [8/9]
<b>9.26.2.19 COUNTERS_TEMPLATE()</b> [9/9]
9.26.2.20 data()

9.26.2.21 desc()	. 281
9.26.2.22 for()	. 281
<b>9.26.2.23 hasher()</b> [1/2]	. 282
<b>9.26.2.24 hasher()</b> [2/2]	. 282
9.26.2.25 hasher_fun() [1/2]	. 282
9.26.2.26 hasher_fun() [2/2]	. 282
9.26.2.27 if() [1/3]	. 282
<b>9.26.2.28 if()</b> [2/3]	. 282
<b>9.26.2.29 if()</b> [3/3]	. 283
<b>9.26.2.30 init_fun()</b> [1/3]	. 283
<b>9.26.2.31 init_fun()</b> [2/3]	. 283
<b>9.26.2.32 init_fun()</b> [3/3]	. 283
9.26.2.33 name()	. 283
9.26.3 Variable Documentation	. 283
9.26.3.1 add_dims	. 283
9.26.3.2 count_fun	. 284
9.26.3.3 counter	. 284
9.26.3.4 counter	. 284
9.26.3.5 data	. 284
9.26.3.6 desc	. 285
9.26.3.7 fun	. 285
9.26.3.8 fun	. 285
9.26.3.9 hasher_fun	. 285
9.26.3.10 i	. 286
9.26.3.11 init_fun	. 286
9.26.3.12 j	. 286
9.26.3.13 name	. 286
9.26.3.14 noexcept	. 286
9.26.3.15 res	. 287
9.26.3.16 return	. 287
9.27 include/barry/counters/network-css.hpp File Reference	. 287
9.27.1 Macro Definition Documentation	. 288
9.27.1.1 CSS_APPEND	. 289
9.27.1.2 CSS_CASE_ELSE	. 289
9.27.1.3 CSS_CASE_PERCEIVED	. 289
9.27.1.4 CSS_CASE_TRUTH	. 289
9.27.1.5 CSS_CHECK_SIZE	. 289
9.27.1.6 CSS_CHECK_SIZE_INIT	. 290
9.27.1.7 CSS_MATCH_TYPE	. 290
9.27.1.8 CSS_NET_COUNTER_LAMBDA_INIT	. 290
9.27.1.9 CSS_PERCEIVED_CELLS	. 290
9.27.1.10 CSS_SIZE	. 291

9.27.1.11 CSS_TRUE_CELLS
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	303
9.31.1 Function Documentation	304
9.31.1.1 likelihood_()	304
9.31.1.2 update_normalizing_constant()	304
9.32 include/barry/models/defm.hpp File Reference	304
9.33 include/barry/models/defm/counters.hpp File Reference	305
9.33.1 Macro Definition Documentation	306
9.33.1.1 DEFM_COUNTER	306
9.33.1.2 DEFM_COUNTER_LAMBDA	307
9.33.1.3 DEFM_RULE	307
9.33.1.4 DEFM_RULE_LAMBDA	307
9.33.1.5 DEFM_RULEDYN_LAMBDA	308
9.33.1.6 UNI_SUB	308
9.34 include/barry/models/geese/counters.hpp File Reference	308
9.35 include/barry/models/defm/defm-bones.hpp File Reference	310
9.36 include/barry/models/defm/defm-meat.hpp File Reference	311
9.36.1 Macro Definition Documentation	311
9.36.1.1 DEFM_LOOP_ARRAYS	311
9.36.1.2 DEFM_RANGES	311
9.36.2 Function Documentation	312
9.36.2.1 keygen_defm()	312
9.37 include/barry/models/defm/defm-types.hpp File Reference	312
9.37.1 Typedef Documentation	313
9.37.1.1 DEFMArray	313
9.38 include/barry/models/defm/formula.hpp File Reference	313
9.38.1 Function Documentation	313
9.38.1.1 defm_motif_parser()	314
9.39 include/barry/models/geese.hpp File Reference	315
9.40 include/barry/models/geese/flock-bones.hpp File Reference	315
9.41 include/barry/models/geese/flock-meat.hpp File Reference	316
9.42 include/barry/models/geese/geese-bones.hpp File Reference	316
9.42.1 Macro Definition Documentation	317
9.42.1.1 INITIALIZED	317
9.42.2 Function Documentation	317
9.42.2.1 keygen_full()	317
9.42.2.2 RULE_FUNCTION()	317
9.42.2.3 vec_diff()	318
9.42.2.4 vector_caster()	318
9.43 include/barry/models/geese/geese-meat-constructors.hpp File Reference	318
9.44 include/barry/models/geese/geese-meat-likelihood.hpp File Reference	319
9.45 include/barry/models/geese/geese-meat-likelihood_exhaust.hpp File Reference	320
9.46 include/barry/models/geese/geese-meat-predict.hpp File Reference	320

9.47 include/barry/models/geese/geese-meat-predict_exhaust.hpp File Reference	321
9.48 include/barry/models/geese/geese-meat-predict_sim.hpp File Reference	321
9.49 include/barry/models/geese/geese-meat-simulate.hpp File Reference	322
9.50 include/barry/models/geese/geese-meat.hpp File Reference	322
9.51 include/barry/models/geese/geese-node-bones.hpp File Reference	323
9.52 include/barry/models/geese/geese-types.hpp File Reference	323
9.52.1 Macro Definition Documentation	324
9.52.1.1 POS	
9.52.2 Typedef Documentation	324
9.52.2.1 PhyloArray	324
9.52.2.2 PhyloCounter	325
9.52.2.3 PhyloCounters	325
9.52.2.4 PhyloModel	325
9.52.2.5 PhyloPowerSet	325
9.52.2.6 PhyloRule	325
9.52.2.7 PhyloRuleData	325
9.52.2.8 PhyloRuleDyn	326
9.52.2.9 PhyloRules	326
9.52.2.10 PhyloRulesDyn	326
9.52.2.11 PhyloStatsCounter	326
9.52.2.12 PhyloSupport	326
9.53 include/barry/powerset-bones.hpp File Reference	326
9.54 include/barry/powerset-meat.hpp File Reference	327
9.55 include/barry/progress.hpp File Reference	327
9.55.1 Macro Definition Documentation	328
9.55.1.1 BARRY_PROGRESS_BAR_WIDTH	328
9.56 include/barry/rules-bones.hpp File Reference	328
9.56.1 Function Documentation	329
9.56.1.1 rule_fun_default()	329
9.57 include/barry/rules-meat.hpp File Reference	329
9.58 include/barry/statscounter-bones.hpp File Reference	329
9.59 include/barry/statscounter-meat.hpp File Reference	330
9.59.1 Macro Definition Documentation	331
9.59.1.1 STATSCOUNTER_TEMPLATE	331
9.59.1.2 STATSCOUNTER_TEMPLATE_ARGS	331
9.59.1.3 STATSCOUNTER_TYPE	331
9.59.2 Function Documentation	331
9.59.2.1 clear()	331
9.59.2.2 for()	332
9.59.2.3 resize()	332
<b>9.59.2.4 STATSCOUNTER_TEMPLATE()</b> [1/9]	332
9.59.2.5 STATSCOUNTER_TEMPLATE() [2/9]	332

9.62.2 Function Documentation	
9.62.1.7 Rule_fun_type	340
9.62.1.6 Row_type	340
9.62.1.5 MapVec_type	
9.62.1.4 Hasher_fun_type	339
9.62.1.3 Counts_type	339
9.62.1.2 Counter_fun_type	
9.62.1.1 Col_type	
9.62.1 Typedef Documentation	
9.62 include/barry/typedefs.hpp File Reference	
9.61.1.1 BARRY_SUPPORT_MEAT_HPP	
9.61.1 Macro Definition Documentation	
9.61 include/barry/support-meat.hpp File Reference	
9.60 include/barry/support-bones.hpp File Reference	
9.59.3.9 return	
9.59.3.8 j	
9.59.3.7 f	
9.59.3.5 current_stats	
9.59.3.4 counters	
9.59.3.3 counters	
9.59.3.2 counter_deleted	
9.59.3.1 counter	
9.59.3 Variable Documentation	333
9.59.2.12 STATSCOUNTER_TEMPLATE() [9/9]	333
9.59.2.11 STATSCOUNTER_TEMPLATE() [8/9]	333
<b>9.59.2.10 STATSCOUNTER_TEMPLATE()</b> [7/9]	333
<b>9.59.2.9 STATSCOUNTER_TEMPLATE()</b> [6/9]	333
<b>9.59.2.8 STATSCOUNTER_TEMPLATE()</b> [5/9]	333
9.59.2.7 STATSCOUNTER_TEMPLATE() [4/9]	332
9.59.2.6 STATSCOUNTER_TEMPLATE() [3/9]	332

# **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	199
NetworkData	
Data class for Networks	200
Node	
A single node for the model	202
NodeData	
Data definition for the PhyloArray class	208
PhyloCounterData	209
PhyloRuleDynData	212
PowerSet < Array_Type, Data_Rule_Type >	
Powerset of a binary array	214
Progress	
A simple progress bar	221
Rule < Array_Type, Data_Type >	
Rule for determining if a cell should be included in a sequence	222
Rules < Array_Type, Data_Type >	
Vector of objects of class Rule	225
StatsCounter< Array_Type, Data_Type >	
Count stats for a single Array	229
Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >	
Compute the support of sufficient statistics	233
vecHasher <t></t>	244

# **Chapter 5**

# File Index

# 5.1 File List

Here is a list of all files with brief descriptions:

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

12 File Index

include/barry/statscounter-meat.hpp
include/barry/support-bones.hpp
include/barry/support-meat.hpp
include/barry/typedefs.hpp
include/barry/counters/network-css.hpp
include/barry/counters/network.hpp
include/barry/models/defm.hpp
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:

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

Definition at line 21 of file counters.hpp.

#### 6.1.2.7 MAKE\_DUPL\_VARS

```
#define MAKE_DUPL_VARS( )

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

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

Definition at line 12 of file counters.hpp.

6.1 Counting 17

#### 6.1.2.8 PHYLO\_CHECK\_MISSING

```
#define PHYLO_CHECK_MISSING( )

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

Definition at line 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

#define PHYLO\_RULE\_DYN\_LAMBDA(

```
Value:
   barry::Rule_fun_type<PhyloArray, PhyloRuleDynData> a = \
   [](const PhyloArray & Array, size_t i, size_t j, PhyloRuleDynData & data)
```

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 > ).
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~(~)} \quad [{\tt 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

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

Keeps track of how many pairs of genes preserve pseudostate.

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

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

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

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

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

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

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

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

Total number of neofunctionalization events.

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

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

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

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

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

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

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

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

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

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

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

# 6.5.1 Detailed Description

Counters for phylogenetic modeling.

**Parameters** 

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

# 6.5.2 Function Documentation

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

Function co-opting.

6.5 Phylo counters 49

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

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

This algorithm implements the change statistic.

Definition at line 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 ti, size ti, Cell< Cell Type > &&v, bool check bounds, bool check exists)
- void insert\_cell (size\_t i, size\_t j, Cell\_Type v, bool check\_bounds, bool check\_exists)
- void swap\_cells (size\_t i0, size\_t j0, size\_t i1, size\_t j1, bool check\_bounds=true, int check\_←
   exists=CHECK::BOTH, int \*report=nullptr)
- void toggle cell (size t i, size t j, bool check bounds=true, int check exists=EXISTS::UKNOWN)
- void toggle\_lock (size\_t i, size\_t j, bool check\_bounds=true)

# Column/row wise interchange

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

# **Arithmetic operators**

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

#### **Public Attributes**

· bool visited = false

### **Friends**

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

#### 8.1.1 Detailed Description

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

Baseline class for binary arrays.

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

### **Template Parameters**

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

62 Class Documentation

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

# 8.1.2 Constructor & Destructor Documentation

# 8.1.2.1 BArray() [1/6]

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

Zero-size array.

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

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

Empty array.

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

# 8.1.2.3 BArray() [3/6]

Edgelist with data.

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

Edgelist with no data (simpler)

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

Copy constructor.

# 8.1.2.6 BArray() [6/6]

Move operator.

# 8.1.2.7 $\sim$ BArray()

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

# 8.1.3 Member Function Documentation

# 8.1.3.1 clear()

64 Class Documentation

# 8.1.3.2 col()

# 8.1.3.3 D() [1/2]

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

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

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

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

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

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

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

# 8.1.3.7 default\_val()

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

# 8.1.3.8 flush\_data()

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

#### 8.1.3.9 get\_cell()

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

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

# 8.1.3.12 get\_entries()

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

# Get the edgelist.

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

#### Returns

Entries < Cell\_Type >

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

66 Class Documentation

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

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

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

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

#### 8.1.3.18 is dense()

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

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

#### 8.1.3.19 is\_empty()

# 8.1.3.20 ncol()

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

### 8.1.3.21 nnozero()

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

# 8.1.3.22 nrow()

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

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

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

68 Class Documentation

#### 8.1.3.25 operator\*=()

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

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

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

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

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

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

# 8.1.3.32 operator/=()

# 8.1.3.33 operator=() [1/2]

Move assignment.

# 8.1.3.34 operator=() [2/2]

Assignment constructor.

#### 8.1.3.35 operator==()

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

70 Class Documentation

# 8.1.3.37 print()

#### 8.1.3.38 print\_n()

#### 8.1.3.39 reserve()

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

# 8.1.3.40 resize()

### 8.1.3.41 rm\_cell()

#### 8.1.3.42 row()

# 8.1.3.43 set\_data()

Set the data object.

#### **Parameters**

data_	
delete_ <i>←</i>	
data_	

### 8.1.3.44 swap\_cells()

# 8.1.3.45 swap\_cols()

72 Class Documentation

### 8.1.3.46 swap\_rows()

# 8.1.3.47 toggle\_cell()

#### 8.1.3.48 toggle\_lock()

# 8.1.3.49 transpose()

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

# 8.1.3.50 zero\_col()

#### 8.1.3.51 zero\_row()

#### 8.1.4 Friends And Related Function Documentation

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

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

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

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

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

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

## 8.1.5 Member Data Documentation

## 8.1.5.1 visited

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

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

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

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

• include/barry/barray-bones.hpp

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

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

#### **Public Member Functions**

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

# 8.2.1 Detailed Description

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

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

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

## 8.2.2 Constructor & Destructor Documentation

## 8.2.2.1 BArrayCell()

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

## 8.2.2.2 ~BArrayCell()

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

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

## 8.2.3 Member Function Documentation

## 8.2.3.1 operator Cell\_Type()

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

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

#### 8.2.3.2 operator\*=()

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

#### 8.2.3.3 operator+=()

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

#### 8.2.3.4 operator-=()

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

#### 8.2.3.5 operator/=()

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

#### 8.2.3.6 operator=()

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

#### 8.2.3.7 operator==()

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

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

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

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

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

## **Public Member Functions**

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

## 8.3.1 Detailed Description

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

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

## 8.3.2 Constructor & Destructor Documentation

## 8.3.2.1 BArrayCell\_const()

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

#### 8.3.2.2 ~BArrayCell\_const()

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

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

#### 8.3.3 Member Function Documentation

#### 8.3.3.1 operator Cell\_Type()

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

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

#### 8.3.3.2 operator"!=()

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

## 8.3.3.3 operator<()

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

#### 8.3.3.4 operator<=()

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

#### 8.3.3.5 operator==()

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

#### 8.3.3.6 operator>()

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

# 8.3.3.7 operator>=()

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

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

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

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

Baseline class for binary arrays.

#include <barraydense-bones.hpp>

#### **Public Member Functions**

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

Get the edgelist.

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

#### Constructors

#### **Parameters**

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

• BArrayDense ()

Zero-size array.

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

Edgelist with no data (simpler)

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

Assignment constructor.

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

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

Move assignment.

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

Set the data object.

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

#### Queries

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

#### **Parameters**

i,j	Coordinates
check_bounds	If false avoids checking bounds.

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

## Cell-wise insertion/deletion

#### Parameters

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

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

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

#### Column/row wise interchange

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

#### **Arithmetic operators**

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

## **Public Attributes**

• bool visited = false

#### **Friends**

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

## 8.4.1 Detailed Description

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

Baseline class for binary arrays.

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

## **Template Parameters**

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

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

## 8.4.2 Constructor & Destructor Documentation

## 8.4.2.1 BArrayDense() [1/6]

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

Zero-size array.

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

## 8.4.2.2 BArrayDense() [2/6]

Empty array.

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

# 8.4.2.3 BArrayDense() [3/6]

Edgelist with data.

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

## 8.4.2.4 BArrayDense() [4/6]

Edgelist with no data (simpler)

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

## 8.4.2.5 BArrayDense() [5/6]

Copy constructor.

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

## 8.4.2.6 BArrayDense() [6/6]

Move operator.

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

## 8.4.2.7 ~BArrayDense()

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

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

## 8.4.3 Member Function Documentation

#### 8.4.3.1 clear()

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

Definition at line 905 of file barraydense-meat.hpp.

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

Definition at line 498 of file barraydense-meat.hpp.

## 8.4.3.3 col() [2/2]

Definition at line 484 of file barraydense-meat.hpp.

# 8.4.3.4 colsum()

Definition at line 1008 of file barraydense-meat.hpp.

## 8.4.3.5 D() [1/2]

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

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

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

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

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

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

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

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

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

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

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

# 8.4.3.9 default\_val()

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

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

#### 8.4.3.10 get\_cell()

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

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

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

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

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

## 8.4.3.13 get\_data()

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

Definition at line 998 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 510 of file barraydense-meat.hpp.

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

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

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

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

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

Definition at line 697 of file barraydense-meat.hpp.

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

Definition at line 663 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 539 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 556 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 560 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 552 of file barraydense-meat.hpp.

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

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

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

Definition at line 626 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 576 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 595 of file barraydense-meat.hpp.

# 8.4.3.33 operator/=()

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

Move assignment.

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

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

Assignment constructor.

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

## 8.4.3.36 operator==()

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

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

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

## 8.4.3.38 print()

Definition at line 964 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 955 of file barraydense-meat.hpp.

#### 8.4.3.40 resize()

Definition at line 919 of file barraydense-meat.hpp.

## 8.4.3.41 rm\_cell()

Definition at line 640 of file barraydense-meat.hpp.

# 8.4.3.42 row() [1/2]

Definition at line 470 of file barraydense-meat.hpp.

## 8.4.3.43 row() [2/2]

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

#### 8.4.3.44 rowsum()

Definition at line 1003 of file barraydense-meat.hpp.

## 8.4.3.45 set\_data()

Set the data object.

#### **Parameters**

data_	
delete_⊸	
data_	

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

## 8.4.3.46 swap\_cells()

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

Definition at line 728 of file barraydense-meat.hpp.

## 8.4.3.47 swap\_cols()

Definition at line 812 of file barraydense-meat.hpp.

#### 8.4.3.48 swap\_rows()

Definition at line 784 of file barraydense-meat.hpp.

## 8.4.3.49 toggle\_cell()

Definition at line 765 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 877 of file barraydense-meat.hpp.

## 8.4.3.52 zero\_col()

Definition at line 858 of file barraydense-meat.hpp.

#### 8.4.3.53 zero\_row()

Definition at line 839 of file barraydense-meat.hpp.

#### 8.4.4 Friends And Related Function Documentation

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

#### 8.4.5 Member Data Documentation

#### 8.4.5.1 visited

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

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

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

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

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

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

#include <barraydensecell-bones.hpp>

#### **Public Member Functions**

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

#### **Friends**

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

## 8.5.1 Detailed Description

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

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

## 8.5.2 Constructor & Destructor Documentation

## 8.5.2.1 BArrayDenseCell()

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

## 8.5.2.2 ~BArrayDenseCell()

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

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

## 8.5.3 Member Function Documentation

## 8.5.3.1 operator Cell\_Type()

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

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

#### 8.5.3.2 operator\*=()

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

#### 8.5.3.3 operator+=()

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

#### 8.5.3.4 operator-=()

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

## 8.5.3.5 operator/=()

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

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

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

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

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

#### 8.5.3.8 operator==()

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

#### 8.5.4 Friends And Related Function Documentation

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

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

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

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

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

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

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

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

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

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

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

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

## 8.6.1 Detailed Description

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

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

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

• include/barry/barraydense-bones.hpp

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

#include <barraydensecol-bones.hpp>

#### **Public Member Functions**

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

## **Friends**

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

## 8.7.1 Detailed Description

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

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

## 8.7.2 Constructor & Destructor Documentation

#### 8.7.2.1 BArrayDenseCol()

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

#### 8.7.3 Member Function Documentation

## 8.7.3.1 begin()

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

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

#### 8.7.3.2 end()

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

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

## 8.7.3.3 operator()()

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

## 8.7.3.4 size()

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

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

## 8.7.4 Friends And Related Function Documentation

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

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

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

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

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

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

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

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

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

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

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

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

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

## **Public Member Functions**

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

#### **Friends**

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

## 8.8.1 Detailed Description

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

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

## 8.8.2 Constructor & Destructor Documentation

#### 8.8.2.1 BArrayDenseCol\_const()

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

# 8.8.3 Member Function Documentation

## 8.8.3.1 begin()

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

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

## 8.8.3.2 end()

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

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

#### 8.8.3.3 operator()()

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

#### 8.8.3.4 size()

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

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

#### 8.8.4 Friends And Related Function Documentation

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

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

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

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

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

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

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

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

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

#include <barraydenserow-bones.hpp>

# **Public Member Functions**

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

#### Friends

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

## 8.9.1 Detailed Description

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

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

#### 8.9.2 Constructor & Destructor Documentation

## 8.9.2.1 BArrayDenseRow()

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

## 8.9.3 Member Function Documentation

## 8.9.3.1 begin()

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

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

## 8.9.3.2 end()

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

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

#### 8.9.3.3 operator()()

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

#### 8.9.3.4 size()

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

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

## 8.9.4 Friends And Related Function Documentation

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

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

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

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

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

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

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

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

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

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

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

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

#include <barraydenserow-bones.hpp>

#### **Public Member Functions**

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

#### **Friends**

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

## 8.10.1 Detailed Description

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

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

## 8.10.2 Constructor & Destructor Documentation

#### 8.10.2.1 BArrayDenseRow\_const()

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

## 8.10.3 Member Function Documentation

#### 8.10.3.1 begin()

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

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

## 8.10.3.2 end()

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

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

#### 8.10.3.3 operator()()

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

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

## 8.10.3.4 size()

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

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

## 8.10.4 Friends And Related Function Documentation

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

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

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

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

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

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

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

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

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

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

#### **Public Member Functions**

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

## 8.11.1 Detailed Description

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

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

## 8.11.2 Constructor & Destructor Documentation

## 8.11.2.1 BArrayRow()

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

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

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

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

## 8.11.3 Member Function Documentation

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

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

## 8.11.3.2 operator\*=()

# 8.11.3.3 operator+=()

#### 8.11.3.4 operator-=()

#### 8.11.3.5 operator/=()

#### 8.11.3.6 operator=()

## 8.11.3.7 operator==()

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

• include/barry/barrayrow-bones.hpp

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

#include <barrayrow-bones.hpp>

# **Public Member Functions**

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

## 8.12.1 Detailed Description

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

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

## 8.12.2 Constructor & Destructor Documentation

#### 8.12.2.1 BArrayRow\_const()

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

#### 8.12.2.2 ~BArrayRow\_const()

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

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

#### 8.12.3 Member Function Documentation

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

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

## 8.12.3.2 operator"!=()

#### 8.12.3.3 operator<()

#### 8.12.3.4 operator<=()

#### 8.12.3.5 operator==()

# 8.12.3.6 operator>()

## 8.12.3.7 operator>=()

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

• include/barry/barrayrow-bones.hpp

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

Row or column of a BArray

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

## **Public Member Functions**

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

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

## 8.13.1 Detailed Description

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

Row or column of a BArray

**Template Parameters** 

Cell_Type	
Data_Type	

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

## 8.13.2 Constructor & Destructor Documentation

## 8.13.2.1 BArrayVector()

Construct a new BArrayVector object.

## **Parameters**

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

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

#### 8.13.2.2 ~BArrayVector()

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

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

#### 8.13.3 Member Function Documentation

## 8.13.3.1 begin()

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

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

#### 8.13.3.2 end()

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

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

# 8.13.3.3 is\_col()

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

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

## 8.13.3.4 is\_row()

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

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

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

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

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

## 8.13.3.6 operator \*= ()

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

#### 8.13.3.7 operator+=()

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

#### 8.13.3.8 operator-=()

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

#### 8.13.3.9 operator/=()

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

## 8.13.3.10 operator=()

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

## 8.13.3.11 operator==()

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

#### 8.13.3.12 size()

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

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

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

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

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

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

#### **Public Member Functions**

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

## 8.14.1 Detailed Description

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

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

#### 8.14.2 Constructor & Destructor Documentation

# 8.14.2.1 BArrayVector\_const()

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

#### 8.14.2.2 ~BArrayVector\_const()

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

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

## 8.14.3 Member Function Documentation

## 8.14.3.1 begin()

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

## 8.14.3.2 end()

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

## 8.14.3.3 is\_col()

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

## 8.14.3.4 is\_row()

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

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

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

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

#### 8.14.3.6 operator"!=()

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

## 8.14.3.7 operator<()

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

## 8.14.3.8 operator<=()

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

# 8.14.3.9 operator==()

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

## 8.14.3.10 operator>()

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

#### 8.14.3.11 operator>=()

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

## 8.14.3.12 size()

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

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

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

# 8.15 Cell< Cell\_Type > Class Template Reference

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

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

#### **Public Member Functions**

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

# **Public Attributes**

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

# 8.15.1 Detailed Description

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

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

· value: the content

· visited: boolean (just a convenient)

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

#### 8.15.2 Constructor & Destructor Documentation

## 8.15.2.1 Cell() [1/7]

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

# 8.15.2.2 Cell() [2/7]

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

## 8.15.2.3 $\sim$ Cell()

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

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

## 8.15.2.4 Cell() [3/7]

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

## 8.15.2.5 Cell() [4/7]

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

## 8.15.2.6 Cell() [5/7]

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

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

## 8.15.2.7 Cell() [6/7]

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

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

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

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

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

## 8.15.3 Member Function Documentation

## 8.15.3.1 add() [1/4]

## 8.15.3.2 add() [2/4]

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

## 8.15.3.3 add() [3/4]

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

Definition at line 52 of file cell-meat.hpp.

## 8.15.3.4 add() [4/4]

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

## 8.15.3.5 operator Cell\_Type()

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

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

# 8.15.3.6 operator"!=()

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

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

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

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

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

## 8.15.3.9 operator==()

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

## 8.15.4 Member Data Documentation

## 8.15.4.1 active

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

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

## 8.15.4.2 value

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

Definition at line 12 of file cell-bones.hpp.

#### 8.15.4.3 visited

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

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

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

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

# 8.16 Cell\_const< Cell\_Type > Class Template Reference

## 8.16.1 Detailed Description

```
template<typename Cell_Type> class Cell_const< Cell_Type>
```

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

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

• include/barry/barray-meat.hpp

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

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

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



## **Public Member Functions**

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

## **Public Attributes**

- · size t current row
- size\_t current\_col
- Row\_type< Cell\_Type >::const\_iterator iter
- const BArray
   Cell\_Type, Data\_Type > \* Array

# 8.17.1 Detailed Description

```
\label{template} \begin{tabular}{ll} template < typename Cell_Type, typename Data_Type > \\ class ConstBArrayRowlter < Cell_Type, Data_Type > \\ \end{tabular}
```

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

## 8.17.2 Constructor & Destructor Documentation

# 8.17.2.1 ConstBArrayRowlter()

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

#### 8.17.2.2 ∼ConstBArrayRowlter()

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

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

#### 8.17.3 Member Data Documentation

#### 8.17.3.1 Array

```
template<typename Cell_Type , typename Data_Type >
const BArray<Cell_Type,Data_Type>* ConstBArrayRowIter< Cell_Type, Data_Type >::Array
```

Definition at line 15 of file barray-iterator.hpp.

#### 8.17.3.2 current\_col

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

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

#### 8.17.3.3 current\_row

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

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

#### 8.17.3.4 iter

```
template<typename Cell_Type , typename Data_Type >
Row_type<Cell_Type>::const_iterator ConstBArrayRowIter< Cell_Type, Data_Type >::iter
```

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

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

• include/barry/barray-iterator.hpp

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

A counter function based on change statistics.

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

## **Public Member Functions**

- ∼Counter ()
- double count (Array\_Type &Array, size\_t i, size\_t j)
- double init (Array\_Type &Array, size\_t i, size\_t j)
- std::string get\_name () const
- std::string get\_description () const

Creator passing a counter and an initializer

#### **Parameters**

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

- Counter ()
- Counter\_fun\_type
   Array\_Type, Data\_Type > count\_fun\_, Counter\_fun\_type
   Array\_Type, Data\_Type > init\_fun\_, Hasher\_fun\_type
   Array\_Type, Data\_Type > hasher\_fun\_, Data\_Type data\_, std::string name\_="", std::string desc\_="")
- Counter (const Counter < Array\_Type, Data\_Type > &counter\_)
   Copy constructor.
- Counter (Counter < Array\_Type, Data\_Type > &&counter\_) noexcept

Move constructor

- Counter< Array\_Type, Data\_Type > operator= (const Counter< Array\_Type, Data\_Type > &counter\_)
   Copy assignment.
- Counter< Array\_Type, Data\_Type > & operator= (Counter< Array\_Type, Data\_Type > &&counter\_)
   noexcept

Move assignment.

void set\_hasher (Hasher\_fun\_type< Array\_Type, Data\_Type > fun)

Get and set the hasher function.

Hasher\_fun\_type< Array\_Type, Data\_Type > get\_hasher ()

## **Public Attributes**

- Counter\_fun\_type
   Array\_Type, Data\_Type > count\_fun
- Counter\_fun\_type
   Array\_Type, Data\_Type > init\_fun
- Hasher\_fun\_type
   Array\_Type, Data\_Type > hasher\_fun
- Data\_Type data
- std::string name = ""
- std::string desc = ""

# 8.18.1 Detailed Description

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

A counter function based on change statistics.

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

Definition at line 35 of file counters-bones.hpp.

## 8.18.2 Constructor & Destructor Documentation

## 8.18.2.1 Counter() [1/4]

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

Definition at line 57 of file counters-bones.hpp.

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

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

# 8.18.2.3 Counter() [3/4]

Copy constructor.

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

Move constructor.

#### 8.18.2.5 ∼Counter()

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

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

#### 8.18.3 Member Function Documentation

## 8.18.3.1 count()

## 8.18.3.2 get\_description()

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

#### 8.18.3.3 get\_hasher()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Hasher_fun_type<Array_Type, Data_Type> Counter< Array_Type, Data_Type >::get_hasher ( )
```

#### 8.18.3.4 get\_name()

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

## 8.18.3.5 init()

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

Copy assignment.

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

Move assignment.

## 8.18.3.8 set\_hasher()

Get and set the hasher function.

The hasher function is used to characterize the support of the array. This way, if possible, the support enumeration is recycled.

**Parameters** 

fun

## 8.18.4 Member Data Documentation

## 8.18.4.1 count\_fun

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

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

#### 8.18.4.2 data

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

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

#### 8.18.4.3 desc

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

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

#### 8.18.4.4 hasher\_fun

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

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

## 8.18.4.5 init\_fun

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

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

## 8.18.4.6 name

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

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

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

• include/barry/counters-bones.hpp

# 8.19 Counters < Array\_Type, Data\_Type > Class Template Reference

Vector of counters.

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

#### **Public Member Functions**

- Counters ()
- ∼Counters ()
- Counters (const Counters < Array\_Type, Data\_Type > &counter\_)

Copy constructor.

Counters (Counters < Array\_Type, Data\_Type > &&counters\_) noexcept

Move constructor.

Counters < Array\_Type, Data\_Type > operator= (const Counters < Array\_Type, Data\_Type > &counter\_)
 Copy assignment constructor.

Counters< Array\_Type, Data\_Type > & operator= (Counters< Array\_Type, Data\_Type > &&counter\_)
 noexcept

Move assignment constructor.

Counter< Array\_Type, Data\_Type > & operator[] (size\_t idx)

Returns a pointer to a particular counter.

• std::size\_t size () const noexcept

Number of counters in the set.

- void add\_counter (Counter< Array\_Type, Data\_Type > counter)
- std::vector< std::string > get\_names () const
- std::vector< std::string > get\_descriptions () const
- std::vector< double > gen\_hash (const Array\_Type &array, bool add\_dims=true)

Generates a hash for the given array according to the counters.

void add\_hash (Hasher\_fun\_type< Array\_Type, Data\_Type > fun\_)

## 8.19.1 Detailed Description

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

Vector of counters.

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

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

#### 8.19.2 Constructor & Destructor Documentation

## 8.19.2.1 Counters() [1/3]

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

## 8.19.2.2 ∼Counters()

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

Definition at line 120 of file counters-bones.hpp.

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

Copy constructor.

## **Parameters**



## 8.19.2.4 Counters() [3/3]

Move constructor.

#### **Parameters**

counters⊷

## 8.19.3 Member Function Documentation

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

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

#### 8.19.3.3 add hash()

## 8.19.3.4 gen\_hash()

Generates a hash for the given array according to the counters.

#### **Parameters**

array	
add_dims	When true (default) the dimmension of the array will be added to the hash.

## Returns

std::vector< double > That can be hashed later.

#### 8.19.3.5 get\_descriptions()

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

# 8.19.3.6 get\_names()

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

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

Copy assignment constructor.

#### **Parameters**



#### Returns

Counters<Array\_Type,Data\_Type>

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

Move assignment constructor.

## **Parameters**



#### Returns

```
Counters<Array_Type,Data_Type>&
```

## 8.19.3.9 operator[]()

Returns a pointer to a particular counter.

#### **Parameters**

```
idx Id of the counter
```

#### Returns

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

## 8.19.3.10 size()

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

Number of counters in the set.

## Returns

size\_t

Definition at line 164 of file counters-bones.hpp.

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

• include/barry/counters-bones.hpp

8.20 DEFM Class Reference 139

# 8.20 DEFM Class Reference

#include <defm-bones.hpp>

Inheritance diagram for DEFM:



Collaboration diagram for DEFM:



## **Public Member Functions**

- DEFM (int \*id, int \*y, double \*x, size\_t id\_length, size\_t y\_ncol, size\_t x\_ncol, size\_t m\_order, bool copy\_
   data=true, bool column\_major=true)
- DEFMModel & get\_model ()
- void init ()
- void simulate (std::vector< double > par, int \*y\_out)
- size\_t get\_n\_y () const
- size\_t get\_n\_obs () const
- size\_t get\_n\_covars () const
- size\_t get\_m\_order () const
- size\_t get\_n\_rows () const
- const int \* get\_Y () const
- const int \* get\_ID () const
- const double \* get\_X () const
- barry::FreqTable< int > motif\_census (std::vector< size\_t > idx)

```
std::vector< double > logodds (const std::vector< double > &par, size_t i, size_t j)
void set_names (std::vector< std::string > Y_names_, std::vector< std::string > X_names_)
const std::vector< std::string > & get_Y_names () const
const std::vector< std::string > & get_X_names () const
void print () const
std::vector< bool > is_motif ()
bool get_column_major () const noexcept
```

## 8.20.1 Detailed Description

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

## 8.20.2 Constructor & Destructor Documentation

#### 8.20.2.1 DEFM()

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

Definition at line 105 of file defm-meat.hpp.

## 8.20.3 Member Function Documentation

```
8.20.3.1 get_column_major()
```

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

Definition at line 449 of file defm-meat.hpp.

# 8.20.3.2 get\_ID()

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

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

8.20 DEFM Class Reference 141

## 8.20.3.3 get\_m\_order()

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

Definition at line 295 of file defm-meat.hpp.

## 8.20.3.4 get\_model()

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

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

## 8.20.3.5 get\_n\_covars()

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

Definition at line 290 of file defm-meat.hpp.

## 8.20.3.6 get\_n\_obs()

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

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

## 8.20.3.7 get\_n\_rows()

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

Definition at line 300 of file defm-meat.hpp.

# 8.20.3.8 get\_n\_y()

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

Definition at line 280 of file defm-meat.hpp.

## 8.20.3.9 get\_X()

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

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

## 8.20.3.10 get\_X\_names()

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

Definition at line 422 of file defm-meat.hpp.

## 8.20.3.11 get\_Y()

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

Definition at line 305 of file defm-meat.hpp.

# 8.20.3.12 get\_Y\_names()

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

Definition at line 418 of file defm-meat.hpp.

## 8.20.3.13 init()

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

Definition at line 215 of file defm-meat.hpp.

# 8.20.3.14 is\_motif()

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

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

8.20 DEFM Class Reference 143

#### 8.20.3.15 logodds()

Definition at line 359 of file defm-meat.hpp.

## 8.20.3.16 motif\_census()

```
barry::FreqTable< int > DEFM::motif_census (  std::vector < size_t > \mathit{idx} ) \quad [inline]
```

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

#### 8.20.3.17 print()

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

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

## 8.20.3.18 set\_names()

```
void DEFM::set_names ( std::vector < std::string > \textit{Y}\_names\_, \\ std::vector < std::string > \textit{X}\_names\_ ) \quad [inline]
```

Definition at line 401 of file defm-meat.hpp.

## 8.20.3.19 simulate()

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

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

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

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

## 8.21 DEFMCounterData Class Reference

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

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

#### **Public Member Functions**

- DEFMCounterData ()
- DEFMCounterData (const std::vector< size\_t > indices\_, const std::vector< double > numbers\_, const std::vector< bool > logical\_, bool is\_motif\_=true)
- size\_t idx (size\_t i) const
- double num (size\_t i) const
- · bool is true (size ti) const
- ∼DEFMCounterData ()

## **Public Attributes**

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

If false, then is a logit intercept.

## 8.21.1 Detailed Description

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

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

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

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

## 8.22 DEFMData Class Reference

Data class for **DEFM** arrays.

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

## **Public Member Functions**

• DEFMData ()

Vector indicating which covariates are included in the model.

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

Constructor.

double operator() (size t i, size t j) const

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

- double at (size\_t i, size\_t j) const
- size\_t ncol () const
- size\_t nrow () const
- · void print () const
- ∼DEFMData ()

## **Public Attributes**

- DEFMArray \* array
- const double \* covariates

Vector of covariates (complete vector)

size\_t obs\_start

Index of the observation in the data.

size\_t X\_ncol

Number of columns in the array of covariates.

size\_t X\_nrow

Number of rows in the array of covariates.

- std::vector< size\_t > covar\_sort
- std::vector< size\_t > covar\_used

Value where the sorting of the covariates is stored.

#### 8.22.1 Detailed Description

Data class for **DEFM** arrays.

This holds information pointing to the data array, including information regarding the number of observations, the time slices of the observation, and the number of covariates in the data.

Definition at line 16 of file defm-types.hpp.

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

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

# 8.23 DEFMRuleData Class Reference

#include <defm-types.hpp>

Inheritance diagram for DEFMRuleData:



#### **Public Member Functions**

- double num (size\_t i) const
- size\_t idx (size\_t i) const
- bool is\_true (size\_t i) const
- DEFMRuleData ()
- DEFMRuleData (std::vector< double > numbers\_, std::vector< size\_t > indices\_, std::vector< bool > logical\_)
- DEFMRuleData (std::vector< double > numbers\_, std::vector< size\_t > indices\_)

## **Public Attributes**

- std::vector< double > numbers
- std::vector< size\_t > indices
- std::vector< bool > logical
- bool init = false

# 8.23.1 Detailed Description

Definition at line 89 of file defm-types.hpp.

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

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

# 8.24 DEFMRuleDynData Class Reference

#include <defm-types.hpp>

Inheritance diagram for DEFMRuleDynData:



Collaboration diagram for DEFMRuleDynData:



# **Public Member Functions**

- DEFMRuleDynData (const std::vector< double > \*counts\_, std::vector< double > numbers\_={}, std::vector< size\_t > indices\_={}, std::vector< bool > logical\_={})
- ∼DEFMRuleDynData ()

## **Public Attributes**

const std::vector< double > \* counts

# 8.24.1 Detailed Description

Definition at line 152 of file defm-types.hpp.

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

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

# 8.25 Entries < Cell\_Type > Class Template Reference

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

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

#### **Public Member Functions**

- Entries ()
- Entries (size\_t n)
- ∼Entries ()
- void resize (size\_t n)

# **Public Attributes**

```
• std::vector< size_t > source
```

- std::vector< size\_t > target
- std::vector< Cell\_Type > val

## 8.25.1 Detailed Description

```
\label{lem:continuous} \begin{split} \text{template} &< \text{typename Cell\_Type} > \\ \text{class Entries} &< \text{Cell\_Type} > \end{split}
```

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

**Template Parameters** 

```
Cell_Type Any type
```

Definition at line 78 of file typedefs.hpp.

## 8.25.2 Constructor & Destructor Documentation

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

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

Definition at line 84 of file typedefs.hpp.

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

Definition at line 85 of file typedefs.hpp.

# 8.25.2.3 $\sim$ Entries()

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

Definition at line 92 of file typedefs.hpp.

## 8.25.3 Member Function Documentation

## 8.25.3.1 resize()

Definition at line 94 of file typedefs.hpp.

## 8.25.4 Member Data Documentation

#### 8.25.4.1 source

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

Definition at line 80 of file typedefs.hpp.

# 8.25.4.2 target

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

Definition at line 81 of file typedefs.hpp.

#### 8.25.4.3 val

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

Definition at line 82 of file typedefs.hpp.

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

• include/barry/typedefs.hpp

## 8.26 Flock Class Reference

A Flock is a group of Geese.

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

#### **Public Member Functions**

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

Add a tree to the flock.

• void set seed (const size t &s)

Set the seed of the model.

- void init (size\_t bar\_width=BARRY\_PROGRESS\_BAR\_WIDTH)
- PhyloCounters \* get\_counters ()
- PhyloSupport \* get support fun ()
- std::vector< 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 225 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 168 of file flock-meat.hpp.

## 8.26.3.11 nleafs()

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

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

## 8.26.3.12 nnodes()

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

Definition at line 182 of file flock-meat.hpp.

# 8.26.3.13 nterms()

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

Definition at line 210 of file flock-meat.hpp.

## 8.26.3.14 ntrees()

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

Definition at line 175 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 303 of file flock-meat.hpp.

# 8.26.3.16 parse\_polytomies()

Check polytomies and return the largest.

Definition at line 232 of file flock-meat.hpp.

## 8.26.3.17 print()

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

Definition at line 259 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 218 of file flock-meat.hpp.

# 8.26.4 Member Data Documentation

# 8.26.4.1 dat

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

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

# 8.26.4.2 initialized

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

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

#### 8.26.4.3 model

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

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

# 8.26.4.4 nfunctions

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

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

#### 8.26.4.5 rengine

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

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

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

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

# 8.27 FreqTable < T > Class Template Reference

Frequency table of vectors.

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

#### **Public Member Functions**

```
• FreqTable ()
```

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

Number of unique elements in the table. (.

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

# 8.27.1 Detailed Description

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

Frequency table of vectors.

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

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

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

Definition at line 22 of file freqtable.hpp.

# 8.27.2 Constructor & Destructor Documentation

# 8.27.2.1 FreqTable()

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

Definition at line 34 of file freqtable.hpp.

## 8.27.2.2 ∼FreqTable()

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

Definition at line 35 of file freqtable.hpp.

## 8.27.3 Member Function Documentation

# 8.27.3.1 add()

Definition at line 59 of file freqtable.hpp.

# 8.27.3.2 as\_vector()

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

Definition at line 139 of file freqtable.hpp.

#### 8.27.3.3 clear()

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

Definition at line 168 of file freqtable.hpp.

## 8.27.3.4 get\_data()

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

Definition at line 40 of file freqtable.hpp.

## 8.27.3.5 get\_index()

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

Definition at line 41 of file freqtable.hpp.

## 8.27.3.6 make\_hash()

Definition at line 239 of file freqtable.hpp.

# 8.27.3.7 print()

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

Definition at line 204 of file freqtable.hpp.

#### 8.27.3.8 reserve()

Definition at line 182 of file freqtable.hpp.

#### 8.27.3.9 size()

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

Number of unique elements in the table. (.

Returns

size t

Definition at line 231 of file freqtable.hpp.

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

• include/barry/freqtable.hpp

## 8.28 Geese Class Reference

Annotated Phylo Model.

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

## **Public Member Functions**

- ∼Geese ()
- void init (size\_t bar\_width=BARRY\_PROGRESS\_BAR\_WIDTH)
- void inherit\_support (const Geese &model\_, bool delete\_support\_=false)
- void calc\_sequence (Node \*n=nullptr)
- void calc reduced sequence ()
- double likelihood (const std::vector< double > &par, bool as\_log=false, bool use\_reduced\_sequence=true, size\_t ncores=1u)
- double likelihood\_exhaust (const std::vector< double > &par)
- std::vector< double > get\_probabilities () const
- void set\_seed (const size\_t &s)
- std::vector< std::vector< size\_t >> simulate (const std::vector< double > &par)
- std::vector< std::vector< double >> observed\_counts ()
- void print\_observed\_counts ()
- void print () const

8.28 Geese Class Reference 161

Prints information about the GEESE.

- void print\_nodes () const
- void init node (Node &n)
- void update annotations (size t nodeid, std::vector< size t > newann)
- std::vector< std::vector< bool >> get\_states () const

Powerset of a gene's possible states.

std::vector< size\_t > get\_annotated\_nodes () const

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

std::vector< size t > get annotations () const

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

## Construct a new Geese object

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

#### **Parameters**

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

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

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

#### Information about the model

#### **Parameters**

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

size\_t nfuns () const noexcept

Number of functions analyzed.

size\_t nnodes () const noexcept

Number of nodes (interior + leaf)

• size\_t nleafs () const noexcept

Number of leaf.

• size t nterms () const

Number of terms included.

• size\_t support\_size () const noexcept

Number of unique sets of sufficient stats.

std::vector< size\_t > nannotations () const noexcept

Number of annotations.

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

Names of the terms in the model.

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

Check polytomies and return the largest.

#### Geese prediction

Calculate the conditional probability

#### **Parameters**

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

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

#### Returns

std::vector< double > Returns the posterior probability

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

   annotated=false, size\_t nsims=10000u)

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

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

#### Returns

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

- std::mt19937 \* get rengine ()
- PhyloCounters \* get\_counters ()
- PhyloModel \* get\_model ()
- PhyloSupport \* get support fun ()

## **Public Attributes**

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

8.28 Geese Class Reference 163

```
    std::vector< std::vector< size_t >> > pset_loc
        Locations of columns.
```

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

#### **Static Public Attributes**

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

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

# 8.28.1 Detailed Description

Annotated Phylo Model.

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

Class representing a phylogenetic tree model with annotations.

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

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

The class provides constructors, a destructor, and methods for initializing the model, inheriting support, calculating the sequence, calculating the reduced sequence, calculating the likelihood, calculating the likelihood exhaustively, getting probabilities, setting the seed, simulating trees, parsing polytomies, getting observed counts, printing observed counts, printing information about the GEESE, and making predictions.

See also

Flock

Definition at line 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.

## 8.28.3.11 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()

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

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:

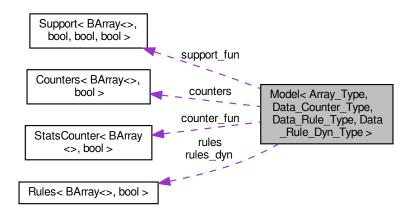
- include/barry/models/geese/geese-bones.hpp
- include/barry/models/geese/geese-meat-constructors.hpp
- include/barry/models/geese/geese-meat-likelihood.hpp
- include/barry/models/geese/geese-meat-likelihood\_exhaust.hpp
- include/barry/models/geese/geese-meat-predict.hpp
- include/barry/models/geese/geese-meat-predict\_exhaust.hpp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

std::vector< std::vector< double >> 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 159 of file model-meat.hpp.

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

Definition at line 193 of file model-meat.hpp.

## 8.29.2.3 Model() [3/3]

Definition at line 231 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.

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

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

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

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

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

Definition at line 349 of file model-meat.hpp.

#### 8.29.3.4 add\_hasher()

Definition at line 384 of file model-meat.hpp.

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

Definition at line 395 of file model-meat.hpp.

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

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

Definition at line 423 of file model-meat.hpp.

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

Definition at line 432 of file model-meat.hpp.

#### 8.29.3.9 colnames()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< std::string > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_\cup Dyn_Type >::colnames [inline]
```

Definition at line 1159 of file model-meat.hpp.

#### 8.29.3.10 conditional prob()

Conditional probability ("Gibbs sampler")

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

#### **Parameters**

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

#### Returns

double The conditional probability

Definition at line 1424 of file model-meat.hpp.

## 8.29.3.11 gen\_key()

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

## 8.29.3.12 get\_arrays2support()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type > std::vector< size_t > * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn \leftarrow _Type >::get_arrays2support [inline]
```

Definition at line 1495 of file model-meat.hpp.

### 8.29.3.13 get\_counters()

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

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

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

Definition at line 1462 of file model-meat.hpp.

#### 8.29.3.14 get\_norm\_const()

Definition at line 949 of file model-meat.hpp.

#### 8.29.3.15 get\_pset()

Definition at line 986 of file model-meat.hpp.

## 8.29.3.16 get\_pset\_arrays()

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

Definition at line 1501 of file model-meat.hpp.

## 8.29.3.17 get\_pset\_probs()

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

Definition at line 1511 of file model-meat.hpp.

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

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

Statistics of the support(s)

Definition at line 1506 of file model-meat.hpp.

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

Definition at line 999 of file model-meat.hpp.

## 8.29.3.20 get\_rengine()

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

Definition at line 1457 of file model-meat.hpp.

## 8.29.3.21 get\_rules()

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

Definition at line 1467 of file model-meat.hpp.

#### 8.29.3.22 get\_rules\_dyn()

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

Definition at line 1472 of file model-meat.hpp.

#### 8.29.3.23 get\_stats\_support()

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

Definition at line 1489 of file model-meat.hpp.

#### 8.29.3.24 get\_stats\_target()

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

Raw pointers to the support and target statistics.

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

Definition at line 1483 of file model-meat.hpp.

#### 8.29.3.25 get support fun()

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

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

Definition at line 1478 of file model-meat.hpp.

#### 8.29.3.26 likelihood() [1/4]

Definition at line 649 of file model-meat.hpp.

## 8.29.3.27 likelihood() [2/4]

Definition at line 803 of file model-meat.hpp.

# 8.29.3.28 likelihood() [3/4]

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

#### 8.29.3.29 likelihood() [4/4]

Definition at line 735 of file model-meat.hpp.

## 8.29.3.30 likelihood\_total()

Definition at line 877 of file model-meat.hpp.

#### 8.29.3.31 nrules()

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

Definition at line 1130 of file model-meat.hpp.

## 8.29.3.32 nrules\_dyn()

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

Definition at line 1138 of file model-meat.hpp.

# 8.29.3.33 nterms()

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

Definition at line 1119 of file model-meat.hpp.

#### 8.29.3.34 operator=()

Definition at line 275 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 1041 of file model-meat.hpp.

#### 8.29.3.36 print\_stats()

Definition at line 1011 of file model-meat.hpp.

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

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

Definition at line 1255 of file model-meat.hpp.

## 8.29.3.38 sample() [2/2]

Definition at line 1175 of file model-meat.hpp.

#### 8.29.3.39 set counters()

Definition at line 366 of file model-meat.hpp.

## 8.29.3.40 set\_rengine()

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

## 8.29.3.41 set\_rules()

Definition at line 405 of file model-meat.hpp.

## 8.29.3.42 set\_rules\_dyn()

Definition at line 447 of file model-meat.hpp.

## 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	

Definition at line 1516 of file model-meat.hpp.

## 8.29.3.45 size()

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

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

Definition at line 1102 of file model-meat.hpp.

## 8.29.3.46 size\_unique()

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

Definition at line 1110 of file model-meat.hpp.

# 8.29.3.47 store\_psets()

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

Definition at line 325 of file model-meat.hpp.

## 8.29.3.48 support\_size()

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

Definition at line 1146 of file model-meat.hpp.

## 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_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
StatsCounter<Array_Type, Data_Counter_Type> Model< Array_Type, Data_Counter_Type, Data_Rule_←
Type, Data_Rule_Dyn_Type >::counter_fun [protected]
```

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

#### 8.29.4.3 counters

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

Definition at line 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_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
MapVec_type< double, size_t > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_\times
Rule_Dyn_Type >::keys2support [protected]
```

Map of types of arrays to support sets.

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

Definition at line 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_←
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 ()
- NetworkData (std::vector< double > vertex\_attr\_, bool directed\_=true)

Constructor using a single attribute.

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

Constructor using multiple attributes.

∼NetworkData ()

## **Public Attributes**

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

# 8.31.1 Detailed Description

Data class for Networks.

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

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

Definition at line 19 of file network.hpp.

## 8.31.2 Constructor & Destructor Documentation

## 8.31.2.1 NetworkData() [1/3]

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

Definition at line 25 of file network.hpp.

## 8.31.2.2 NetworkData() [2/3]

Constructor using a single attribute.

#### Parameters

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

Definition at line 33 of file network.hpp.

## 8.31.2.3 NetworkData() [3/3]

Constructor using multiple attributes.

#### **Parameters**

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

Definition at line 45 of file network.hpp.

## 8.31.2.4 ~NetworkData()

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

Definition at line 51 of file network.hpp.

# 8.31.3 Member Data Documentation

#### 8.31.3.1 directed

bool NetworkData::directed = true

Definition at line 22 of file network.hpp.

## 8.31.3.2 vertex\_attr

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

Definition at line 23 of file network.hpp.

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

• include/barry/counters/network.hpp

# 8.32 Node Class Reference

A single node for the model.

#include <geese-node-bones.hpp>

Collaboration diagram for Node:



8.32 Node Class Reference 203

#### **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\_t ord

Order in which the node was created.

PhyloArray array

Array of the node.

std::vector< size\_t > annotations

Observed annotations (only defined for Geese)

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

Arrays given all possible states.

Node \* parent = nullptr

Parent node.

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

Offspring nodes.

•  $std::vector < size_t > narray = {}$ 

ID of the array in the model.

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

Induced subtree probabilities.

std::vector< double > probability

The probability of observing each state.

# 8.32.1 Detailed Description

A single node for the model.

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

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

## 8.32.2 Constructor & Destructor Documentation

# 8.32.2.1 Node() [1/5]

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

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

## 8.32.2.2 Node() [2/5]

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

## 8.32.2.3 Node() [3/5]

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

## 8.32.2.4 Node() [4/5]

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

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

## **8.32.2.5** Node() [5/5]

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

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

8.32 Node Class Reference 205

## 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.

# 8.32.4.3 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 Node Class Reference 207

# 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 15 of file geese-types.hpp.

## 8.33.2 Constructor & Destructor Documentation

## 8.33.2.1 NodeData()

Definition at line 35 of file geese-types.hpp.

#### 8.33.3 Member Data Documentation

#### 8.33.3.1 blengths

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

Branch length.

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

## 8.33.3.2 duplication

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

Definition at line 31 of file geese-types.hpp.

#### 8.33.3.3 states

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

State of the parent node.

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

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

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

# 8.34 PhyloCounterData Class Reference

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

## **Public Member Functions**

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

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

# 8.34.1 Detailed Description

Definition at line 45 of file geese-types.hpp.

# 8.34.2 Constructor & Destructor Documentation

# 8.34.2.1 PhyloCounterData() [1/2]

Definition at line 51 of file geese-types.hpp.

## 8.34.2.2 PhyloCounterData() [2/2]

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

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

# 8.34.3 Member Function Documentation

## 8.34.3.1 at()

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

# 8.34.3.2 begin()

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

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

## 8.34.3.3 empty()

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

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

# 8.34.3.4 end()

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

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

## 8.34.3.5 get\_counters()

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

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

#### 8.34.3.6 operator()()

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

# 8.34.3.7 operator[]()

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

# 8.34.3.8 push\_back()

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

## 8.34.3.9 reserve()

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

#### 8.34.3.10 shrink\_to\_fit()

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

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

## 8.34.3.11 size()

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

Definition at line 64 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 74 of file geese-types.hpp.

#### 8.35.2 Constructor & Destructor Documentation

## 8.35.2.1 PhyloRuleDynData()

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

## 8.35.2.2 ~PhyloRuleDynData()

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

Definition at line 96 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 91 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 76 of file geese-types.hpp.

# 8.35.4.2 duplication

size\_t PhyloRuleDynData::duplication

Definition at line 80 of file geese-types.hpp.

#### 8.35.4.3 lb

size\_t PhyloRuleDynData::lb

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

## 8.35.4.4 pos

size\_t PhyloRuleDynData::pos

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

#### 8.35.4.5 ub

size\_t PhyloRuleDynData::ub

Definition at line 79 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_←	When true, the Rule destructor will delete the pointer, if defined.
dat_	

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

### 8.38.1 Detailed Description

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

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

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

#### **Template Parameters**

Array_Type	An object of class BArray.
Data_Type	Any type.

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

## 8.38.2 Constructor & Destructor Documentation

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

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

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

#### 8.38.2.2 Rule() [2/2]

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

#### 8.38.2.3 ∼Rule()

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

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

#### 8.38.3 Member Function Documentation

## 8.38.3.1 D()

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

Read/Write access to the data.

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

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

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

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

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

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

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

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

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

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

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

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

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

#### 8.38.3.6 operator()()

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

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

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

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

Vector of objects of class Rule.

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

#### **Public Member Functions**

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

Check whether a given cell is free or locked.

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

#### Rule adding

#### **Parameters**

```
rule
```

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

## 8.39.1 Detailed Description

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

Vector of objects of class Rule.

#### **Template Parameters**

Array_Type	An object of class BArray
Data_Type	Any type.

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

## 8.39.2 Constructor & Destructor Documentation

## 8.39.2.1 Rules() [1/2]

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

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

#### 8.39.2.2 Rules() [2/2]

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

#### 8.39.2.3 ∼Rules()

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

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

#### 8.39.3 Member Function Documentation

#### 8.39.3.1 add rule() [1/2]

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

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

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

## 8.39.3.3 begin()

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

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

## 8.39.3.4 end()

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

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

#### 8.39.3.5 get\_descriptions()

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

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

#### 8.39.3.6 get\_names()

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

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

## 8.39.3.7 get\_seq()

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

## **Parameters**

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

#### Returns

Nothing.

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

## 8.39.3.8 operator()()

Check whether a given cell is free or locked.

#### **Parameters**

а	A BArray object
i	row position
j	col position

#### Returns

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

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

#### 8.39.3.9 operator=()

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

## 8.39.3.10 size()

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

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

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

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

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

Count stats for a single Array.

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

#### **Public Member Functions**

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

Creator of a StatsCounter

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

Copy constructor.

• StatsCounter ()

Can be created without setting the array.

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

Changes the reference array for the counting.

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

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

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

## 8.40.1 Detailed Description

```
\label{template} \begin{tabular}{ll} template < typename \ Array\_Type, \ typename \ Data\_Type > \\ class \ Stats Counter < Array\_Type, \ Data\_Type > \\ \end{tabular}
```

Count stats for a single Array.

Users can a list of functions that can be used with this. The baseline set of arguments is a pointer to a binary array and a dataset to add the counts to.

Definition at line 14 of file statscounter-bones.hpp.

#### 8.40.2 Constructor & Destructor Documentation

## 8.40.2.1 StatsCounter() [1/3]

Creator of a StatsCounter

#### **Parameters**

Array←	A const pointer to a BArray.

Definition at line 37 of file statscounter-bones.hpp.

## 8.40.2.2 StatsCounter() [2/3]

Copy constructor.

#### **Parameters**

counter

### 8.40.2.3 StatsCounter() [3/3]

```
template<typename Array_Type , typename Data_Type >
StatsCounter< Array_Type, Data_Type >::StatsCounter ( ) [inline]
```

Can be created without setting the array.

Definition at line 59 of file statscounter-bones.hpp.

## 8.40.2.4 $\sim$ StatsCounter()

```
template<typename Array_Type , typename Data_Type >
StatsCounter< Array_Type, Data_Type >::~StatsCounter ( )
```

## 8.40.3 Member Function Documentation

### 8.40.3.1 add\_counter()

#### 8.40.3.2 count\_all()

```
template<typename Array_Type , typename Data_Type >
std::vector< double > StatsCounter< Array_Type, Data_Type >::count_all [inline]
```

Definition at line 99 of file statscounter-meat.hpp.

#### 8.40.3.3 count\_current()

## 8.40.3.4 count\_init()

Counter functions This function recurses through the entries of Array and at each step of adding a new cell it uses the functions to list the statistics.

## 8.40.3.5 get\_counters()

```
template<typename Array_Type , typename Data_Type >
Counters<Array_Type,Data_Type>* StatsCounter< Array_Type, Data_Type >::get_counters ( )
```

### 8.40.3.6 get\_descriptions()

```
template<typename Array_Type , typename Data_Type >
std::vector< std::string > StatsCounter< Array_Type, Data_Type >::get_descriptions ( ) const
```

#### 8.40.3.7 get\_names()

```
template<typename Array_Type , typename Data_Type >
std::vector< std::string > StatsCounter< Array_Type, Data_Type >::get_names ( ) const
```

#### 8.40.3.8 reset array()

Changes the reference array for the counting.

#### **Parameters**

Array←	A pointer to an array of class Array_Type.
1_	

#### 8.40.3.9 set counters()

#### 8.40.3.10 size()

```
template<typename Array_Type , typename Data_Type >
size_t StatsCounter< Array_Type, Data_Type >::size ( ) const [inline]
```

Definition at line 86 of file statscounter-bones.hpp.

The documentation for this class was generated from the following files:

- include/barry/statscounter-bones.hpp
- include/barry/statscounter-meat.hpp

# 8.41 Support < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type > Class Template Reference

Compute the support of sufficient statistics.

```
#include <support-bones.hpp>
```

#### **Public Member Functions**

Support (const Array\_Type &Array\_)

Constructor passing a reference Array.

• Support (size\_t N\_, size\_t M\_)

Constructor specifying the dimensions of the array (empty).

- Support ()
- ∼Support ()
- void init\_support (std::vector< Array\_Type > \*array\_bank=nullptr, std::vector< double > \*stats\_\cup bank=nullptr)
- void calc (std::vector < Array\_Type > \*array\_bank=nullptr, std::vector < double > \*stats\_bank=nullptr, size ←
   \_t max\_num\_elements\_=0u)

Computes the entire support.

- 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  $rule_{dyn}$  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]

Definition at line 478 of file support-meat.hpp.

## 8.41.3.5 add\_rule\_dyn() [2/2]

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_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 \( \to \) _Type >::change_stats
```

Definition at line 82 of file support-bones.hpp.

## 8.41.4.2 coordiantes\_n\_free

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::coordiantes 
n free
```

Definition at line 80 of file support-bones.hpp.

#### 8.41.4.3 coordiantes\_n\_locked

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::coordiantes \cdot _n_locked
```

Definition at line 81 of file support-bones.hpp.

#### 8.41.4.4 coordinates free

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< size_t > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn 
__Type >::coordinates_free
```

Definition at line 78 of file support-bones.hpp.

#### 8.41.4.5 coordinates\_locked

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< size_t > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn 
_Type >::coordinates_locked
```

Definition at line 79 of file support-bones.hpp.

## 8.41.4.6 current\_stats

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< double > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn← _Type >::current_stats
```

Definition at line 77 of file support-bones.hpp.

## 8.41.4.7 delete\_counters

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_← counters = true
```

Definition at line 71 of file support-bones.hpp.

#### 8.41.4.8 delete\_rules

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_← rules = true
```

Definition at line 72 of file support-bones.hpp.

#### 8.41.4.9 delete rules dyn

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_← rules_dyn = true
```

Definition at line 73 of file support-bones.hpp.

#### 8.41.4.10 hashes

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< size_t > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn← _Type >::hashes
```

Definition at line 83 of file support-bones.hpp.

## 8.41.4.11 hashes\_initialized

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< bool > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_↔ Type >::hashes_initialized
```

Definition at line 84 of file support-bones.hpp.

#### 8.41.4.12 M

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename
Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::M
```

Definition at line 70 of file support-bones.hpp.

#### 8.41.4.13 max\_num\_elements

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename
Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::max_num
_elements = BARRY_MAX_NUM_ELEMENTS
```

Definition at line 74 of file support-bones.hpp.

#### 8.41.4.14 N

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename
Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::N
```

Definition at line 70 of file support-bones.hpp.

#### 8.41.4.15 n counters

```
template<typename Array_Type = BArray<bool, bool>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::n_
counters
```

Definition at line 85 of file support-bones.hpp.

The documentation for this class was generated from the following files:

- include/barry/support-bones.hpp
- include/barry/support-meat.hpp

## 8.42 vecHasher < T > Struct Template Reference

#include <typedefs.hpp>

#### **Public Member Functions**

• std::size\_t operator() (std::vector< T > const &dat) const noexcept

## 8.42.1 Detailed Description

```
\label{eq:typename} \begin{array}{l} \text{template}\!<\!\text{typename T}\!>\\ \text{struct vecHasher}\!<\!\text{T}\!> \end{array}
```

Definition at line 105 of file typedefs.hpp.

## 8.42.2 Member Function Documentation

#### 8.42.2.1 operator()()

Definition at line 108 of file typedefs.hpp.

The documentation for this struct was generated from the following file:

• include/barry/typedefs.hpp

## **Chapter 9**

## **File Documentation**

## 9.1 include/barry/barray-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



## **Classes**

class BArray < Cell\_Type, Data\_Type >
 Baseline class for binary arrays.

## 9.2 include/barry/barray-iterator.hpp File Reference

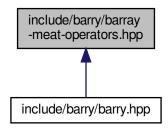
#### Classes

class ConstBArrayRowIter< Cell\_Type, Data\_Type >

248 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

250 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]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator+ ) const &
```

Definition at line 46 of file barray-meat-operators.hpp.

## 9.3.2.4 BARRAY\_TEMPLATE() [4/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator- ) const
```

## 9.3.2.5 BARRAY\_TEMPLATE() [5/6]

Definition at line 75 of file barray-meat-operators.hpp.

## **9.3.2.6 BARRAY\_TEMPLATE()** [6/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator/ ) const &
```

Definition at line 105 of file barray-meat-operators.hpp.

## 9.3.2.7 BARRAY\_TEMPLATE\_ARGS()

```
template BARRAY_TEMPLATE_ARGS ( ) const \&
```

## 9.3.2.8 BARRAY\_TYPE()

```
template const BARRAY_TYPE ( ) &
```

Definition at line 20 of file barray-meat-operators.hpp.

#### 9.3.2.9 for()

```
for ( ) [pure virtual]
```

Definition at line 66 of file barray-meat-operators.hpp.

## 9.3.2.10 operator()()

## 9.3.3 Variable Documentation

252 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.

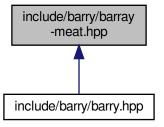
#### 9.3.3.2 this

```
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 >

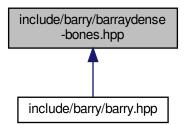
# 9.6 include/barry/barraycell-meat.hpp File Reference

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 &
```

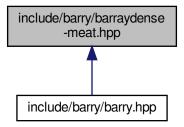
## 9.8.2.6 BDENSE\_TYPE()

```
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

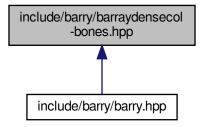
#### 9.11.1.1 POS

```
#define POS(  a, \\ b ) \ (a) \ + \ (b) \ * \ dat -> \mathbb{N}
```

Definition at line 6 of file barraydensecell-meat.hpp.

# 9.12 include/barry/barraydensecol-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Classes**

- class BArrayDenseCol< Cell\_Type, Data\_Type >
- class BArrayDenseCol\_const< Cell\_Type, Data\_Type >

## **Macros**

- #define POS(a, b) (b)\*N + (a)
- #define POS\_N(a, b, c) (b)\*(c) + (a)
- #define ZERO\_CELL static\_cast<Cell\_Type>(0.0)

#### 9.12.1 Macro Definition Documentation

# 9.12.1.1 POS

```
#define POS(  a, \\ b ) (b)*N + (a)
```

Definition at line 4 of file barraydensecol-bones.hpp.

## 9.12.1.2 POS\_N

Definition at line 5 of file barraydensecol-bones.hpp.

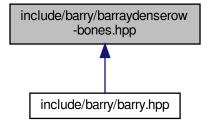
# 9.12.1.3 ZERO\_CELL

```
#define ZERO_CELL static_cast<Cell_Type>(0.0)
```

Definition at line 6 of file barraydensecol-bones.hpp.

# 9.13 include/barry/barraydenserow-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



### Classes

- class BArrayDenseRow
   Cell\_Type, Data\_Type >
- class BArrayDenseRow\_const< Cell\_Type, Data\_Type >

## **Macros**

- #define POS(a, b) (b) \* N + (a)
- #define POS\_N(a, b, c) (b)\*(c) + (a)
- #define ZERO\_CELL static\_cast< Cell\_Type >(0.0)

#### 9.13.1 Macro Definition Documentation

#### 9.13.1.1 POS

```
#define POS(  a, \\ b ) \ (b) \ * \ N \ + \ (a)
```

Definition at line 4 of file barraydenserow-bones.hpp.

#### 9.13.1.2 POS N

Definition at line 5 of file barraydenserow-bones.hpp.

#### 9.13.1.3 ZERO\_CELL

```
#define ZERO_CELL static_cast< Cell_Type >(0.0)
```

Definition at line 6 of file barraydenserow-bones.hpp.

# 9.14 include/barry/barrayrow-bones.hpp File Reference

#### **Classes**

- class BArrayRow
   Cell\_Type, Data\_Type >
- class BArrayRow const< Cell Type, Data Type >

# 9.15 include/barry/barrayrow-meat.hpp File Reference

### **Macros**

- #define BROW\_TYPE() BArrayRow<Cell\_Type, Data\_Type>
- #define BROW\_TEMPLATE\_ARGS() < typename Cell\_Type, typename Data\_Type >
- #define BROW\_TEMPLATE(a, b) template BROW\_TEMPLATE\_ARGS() inline a BROW\_TYPE()::b

#### **Functions**

- BROW\_TEMPLATE (void, operator=)(const BROW\_TYPE() &val)
- BROW\_TEMPLATE (void, operator+=)(const BROW\_TYPE() &val)
- BROW\_TEMPLATE (void, operator-=)(const BROW\_TYPE() &val)
- BROW TEMPLATE (void, operator\*=)(const BROW TYPE() &val)
- BROW\_TEMPLATE (void, operator/=)(const BROW\_TYPE() &val)

# 9.15.1 Macro Definition Documentation

#### 9.15.1.1 BROW\_TEMPLATE

Definition at line 8 of file barrayrow-meat.hpp.

#### 9.15.1.2 BROW\_TEMPLATE\_ARGS

```
#define BROW_TEMPLATE_ARGS() <typename Cell_Type, typename Data_Type>
```

Definition at line 6 of file barrayrow-meat.hpp.

## 9.15.1.3 BROW\_TYPE

```
#define BROW_TYPE( ) BArrayRow<Cell_Type, Data_Type>
```

Definition at line 4 of file barrayrow-meat.hpp.

# 9.15.2 Function Documentation

### 9.15.2.1 BROW\_TEMPLATE() [1/5]

Definition at line 45 of file barrayrow-meat.hpp.

## 9.15.2.2 BROW\_TEMPLATE() [2/5]

Definition at line 25 of file barrayrow-meat.hpp.

## 9.15.2.3 BROW\_TEMPLATE() [3/5]

Definition at line 34 of file barrayrow-meat.hpp.

#### 9.15.2.4 BROW\_TEMPLATE() [4/5]

Definition at line 55 of file barrayrow-meat.hpp.

# 9.15.2.5 BROW\_TEMPLATE() [5/5]

Definition at line 11 of file barrayrow-meat.hpp.

# 9.16 include/barry/barrayvector-bones.hpp File Reference

#### **Classes**

```
    class BArrayVector< Cell_Type, Data_Type >
    Row or column of a BArray
```

class BArrayVector\_const< Cell\_Type, Data\_Type >

# 9.17 include/barry/barrayvector-meat.hpp File Reference

# 9.18 include/barry/barry-configuration.hpp File Reference

This graph shows which files directly or indirectly include this file:



# **Configuration MACROS**

These are mostly related to performance. The definitions follow:

- BARRY\_USE\_UNORDERED\_MAP If specified, then barry is compiled using std::unordered\_map. Otherwise it will use std::map for the arrays.
- BARRY\_USE\_SAFE\_EXP When specified, it will multiply all likelihoods in Model by (1/-100)/(1/-100) so that numerical overflows are avoided.
- BARRY\_USE\_ISFINITE When specified, it will introduce a macro that checks whether the likelihood is finite or not.
- $printf\_barry$  If not specified, will be defined as printf.
- BARRY\_DEBUG\_LEVEL, when defined, will make things verbose.
- #define BARRY\_SAFE\_EXP -100.0
- #define BARRY ISFINITE(a)
- #define BARRY\_CHECK\_SUPPORT(x, maxs)
- #define printf\_barry printf
- #define BARRY\_MAX\_NUM\_ELEMENTS static\_cast< size\_t >(std::numeric\_limits< size\_t >::max() /2u)
- template<typename Ta , typename Tb >
   using Map = std::map< Ta, Tb >

#### 9.18.1 Macro Definition Documentation

#### 9.18.1.1 BARRY CHECK SUPPORT

```
#define BARRY_CHECK_SUPPORT(
          x,
          maxs )
```

Definition at line 47 of file barry-configuration.hpp.

#### 9.18.1.2 BARRY\_ISFINITE

Definition at line 40 of file barry-configuration.hpp.

# 9.18.1.3 BARRY\_MAX\_NUM\_ELEMENTS

```
#define BARRY_MAX_NUM_ELEMENTS static_cast< size_t >(std::numeric_limits< size_t >::max()
/2u)
```

Definition at line 55 of file barry-configuration.hpp.

#### 9.18.1.4 BARRY\_SAFE\_EXP

```
#define BARRY_SAFE_EXP -100.0
```

Definition at line 33 of file barry-configuration.hpp.

# 9.18.1.5 printf\_barry

```
#define printf_barry printf
```

Definition at line 51 of file barry-configuration.hpp.

# 9.18.2 Typedef Documentation

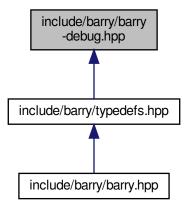
#### 9.18.2.1 Map

```
template<typename Ta , typename Tb >
using Map = std::map<Ta,Tb>
```

Definition at line 27 of file barry-configuration.hpp.

# 9.19 include/barry/barry-debug.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Macros**

• #define BARRY\_DEBUG\_LEVEL 0

## 9.19.1 Macro Definition Documentation

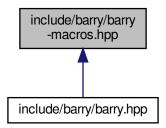
## 9.19.1.1 BARRY\_DEBUG\_LEVEL

#define BARRY\_DEBUG\_LEVEL 0

Definition at line 5 of file barry-debug.hpp.

# 9.20 include/barry/barry-macros.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Macros**

- #define BARRY\_ZERO Cell
   Cell\_Type>(0.0)
- #define BARRY\_ZERO\_DENSE static\_cast<Cell\_Type>(0.0)
- #define BARRY\_ONE Cell<Cell\_Type>(1.0)
- #define BARRY\_ONE\_DENSE static\_cast<Cell\_Type>(1.0)
- #define BARRY\_UNUSED(expr) do { (void)(expr); } while (0);
- #define BARRY\_NCORES\_ARG(default) size\_t ncores default

#### 9.20.1 Macro Definition Documentation

### 9.20.1.1 BARRY\_NCORES\_ARG

Definition at line 15 of file barry-macros.hpp.

## 9.20.1.2 BARRY\_ONE

```
#define BARRY_ONE CellCell_Type>(1.0)
```

Definition at line 7 of file barry-macros.hpp.

## 9.20.1.3 BARRY\_ONE\_DENSE

```
#define BARRY_ONE_DENSE static_cast<Cell_Type>(1.0)
```

Definition at line 8 of file barry-macros.hpp.

#### 9.20.1.4 BARRY\_UNUSED

Definition at line 10 of file barry-macros.hpp.

#### 9.20.1.5 BARRY\_ZERO

```
#define BARRY_ZERO Cell<Cell_Type>(0.0)
```

Definition at line 4 of file barry-macros.hpp.

#### 9.20.1.6 BARRY\_ZERO\_DENSE

```
#define BARRY_ZERO_DENSE static_cast<Cell_Type>(0.0)
```

Definition at line 5 of file barry-macros.hpp.

# 9.21 include/barry/barry.hpp File Reference

```
#include <iostream>
#include <cstdarg>
#include <vector>
#include <unordered_map>
#include <functional>
#include <stdexcept>
#include <cmath>
#include <map>
#include <algorithm>
#include <utility>
#include <random>
#include <climits>
#include <cfloat>
#include <string>
#include <cstdint>
#include <memory>
#include <regex>
```

```
#include <iterator>
#include "typedefs.hpp"
#include "barry-macros.hpp"
#include "freqtable.hpp"
#include "cell-bones.hpp"
#include "cell-meat.hpp"
#include "barray-bones.hpp"
#include "barraycell-bones.hpp"
#include "barray-meat.hpp"
#include "barraycell-meat.hpp"
#include "barray-meat-operators.hpp"
#include "barraydense-bones.hpp"
#include "barraydensecell-bones.hpp"
#include "barraydenserow-bones.hpp"
#include "barraydensecol-bones.hpp"
#include "barraydense-meat.hpp"
#include "barraydensecell-meat.hpp"
#include "barraydense-meat-operators.hpp"
#include "counters-bones.hpp"
#include "counters-meat.hpp"
#include "statscounter-bones.hpp"
#include "statscounter-meat.hpp"
#include "support-bones.hpp"
#include "support-meat.hpp"
#include "powerset-bones.hpp"
#include "powerset-meat.hpp"
#include "model-bones.hpp"
#include "model-meat.hpp"
#include "rules-bones.hpp"
#include "rules-meat.hpp"
#include "counters/network.hpp"
Include dependency graph for barry.hpp:
```

#### **Namespaces**

barry

barry: Your go-to motif accountant

barry::counters

Tree class and Treelterator class.

· barry::counters::network

#### **Macros**

- #define BARRY HPP
- #define BARRY VERSION MAYOR 0
- #define BARRY\_VERSION\_MINOR 1
- #define BARRY\_VERSION BARRY\_VERSION\_MAYOR ## . ## BARRY\_VERSION\_MINOR
- #define COUNTER\_FUNCTION(a)
- #define COUNTER\_LAMBDA(a)
- #define RULE\_FUNCTION(a)
- #define RULE\_LAMBDA(a)

#### 9.21.1 Macro Definition Documentation

#### 9.21.1.1 BARRY HPP

```
#define BARRY_HPP
```

Definition at line 29 of file barry.hpp.

#### 9.21.1.2 BARRY\_VERSION

```
#define BARRY_VERSION BARRY_VERSION_MAYOR ## . ## BARRY_VERSION_MINOR
```

Definition at line 33 of file barry.hpp.

#### 9.21.1.3 BARRY\_VERSION\_MAYOR

```
#define BARRY_VERSION_MAYOR 0
```

Definition at line 31 of file barry.hpp.

# 9.21.1.4 BARRY\_VERSION\_MINOR

```
#define BARRY_VERSION_MINOR 1
```

Definition at line 32 of file barry.hpp.

## 9.21.1.5 COUNTER\_FUNCTION

```
\begin{array}{c} \texttt{\#define COUNTER\_FUNCTION}\,(\\ & a \end{array})
```

#### Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
inline double (a) (const Array_Type & Array, size_t i, size_t j, Data_Type & data)\
```

Definition at line 92 of file barry.hpp.

## 9.21.1.6 COUNTER\_LAMBDA

Definition at line 95 of file barry.hpp.

#### 9.21.1.7 RULE\_FUNCTION

Definition at line 99 of file barry.hpp.

#### 9.21.1.8 RULE LAMBDA

Definition at line 102 of file barry.hpp.

# 9.22 include/barry/cell-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Classes**

class Cell
 Cell\_Type >
 Entries in BArray. For now, it only has two members:

# 9.23 include/barry/cell-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



# 9.24 include/barry/col-bones.hpp File Reference

# 9.25 include/barry/counters-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:

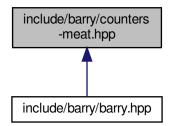


### **Classes**

- class Counter< Array\_Type, Data\_Type >
  - A counter function based on change statistics.
- class Counters
   Array\_Type, Data\_Type >
   Vector of counters.

# 9.26 include/barry/counters-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Macros**

- #define COUNTER\_TYPE() Counter<Array\_Type,Data\_Type>
- #define COUNTER\_TEMPLATE\_ARGS() < typename Array\_Type, typename Data\_Type>
- #define 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
- 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\_
- Data\_Type Counter\_fun\_type
   Array\_Type, Data\_Type > Hasher\_fun\_type
   Array\_Type, Data\_Type > Data\_Type std::string name
- Data\_Type Counter\_fun\_type< Array\_Type, Data\_Type > Hasher\_fun\_type< Array\_Type, Data\_Type > Data\_Type std::string std::string desc\_
- · bool add dims
- · return res
- Data Type fun

#### 9.26.1 Macro Definition Documentation

#### 9.26.1.1 COUNTER\_TEMPLATE

Definition at line 8 of file counters-meat.hpp.

#### 9.26.1.2 COUNTER\_TEMPLATE\_ARGS

```
#define COUNTER_TEMPLATE_ARGS() <typename Array_Type, typename Data_Type>
```

Definition at line 6 of file counters-meat.hpp.

#### 9.26.1.3 COUNTER\_TYPE

```
#define COUNTER_TYPE( ) Counter<Array_Type, Data_Type>
```

Definition at line 4 of file counters-meat.hpp.

#### 9.26.1.4 COUNTERS TEMPLATE

Definition at line 129 of file counters-meat.hpp.

#### 9.26.1.5 COUNTERS\_TEMPLATE\_ARGS

```
#define COUNTERS_TEMPLATE_ARGS() <typename Array_Type, typename Data_Type>
```

Definition at line 127 of file counters-meat.hpp.

### 9.26.1.6 COUNTERS\_TYPE

```
#define COUNTERS_TYPE( ) Counters<Array_Type,Data_Type>
```

Definition at line 125 of file counters-meat.hpp.

## 9.26.1.7 TMP\_HASHER\_CALL

```
#define TMP_HASHER_CALL Hasher_fun_type<Array_Type,Data_Type>
```

Definition at line 115 of file counters-meat.hpp.

#### 9.26.2 Function Documentation

#### 9.26.2.1 count fun()

# 9.26.2.2 COUNTER\_TEMPLATE() [1/9]

```
COUNTER_TEMPLATE (

Counter ) const
```

# 9.26.2.3 COUNTER\_TEMPLATE() [2/9]

# 9.26.2.4 COUNTER\_TEMPLATE() [3/9]

# 9.26.2.5 COUNTER\_TEMPLATE() [4/9]

< Move assignment

# **9.26.2.6 COUNTER\_TEMPLATE()** [5/9]

# 9.26.2.7 **COUNTER\_TEMPLATE()** [6/9]

```
COUNTER_TEMPLATE (
          std::string ,
          get_description ) const
```

Definition at line 107 of file counters-meat.hpp.

## 9.26.2.8 **COUNTER\_TEMPLATE()** [7/9]

```
COUNTER_TEMPLATE (
          std::string ,
          get_name ) const
```

Definition at line 103 of file counters-meat.hpp.

### 9.26.2.9 COUNTER\_TEMPLATE() [8/9]

```
COUNTER_TEMPLATE (

TMP_HASHER_CALL ,

qet_hasher )
```

Definition at line 116 of file counters-meat.hpp.

#### 9.26.2.10 COUNTER\_TEMPLATE() [9/9]

# 9.26.2.11 COUNTERS\_TEMPLATE() [1/9]

```
COUNTERS_TEMPLATE (

Counters )
```

Definition at line 132 of file counters-meat.hpp.

#### 9.26.2.12 COUNTERS\_TEMPLATE() [2/9]

```
COUNTERS_TEMPLATE (

COUNTER_TYPE() & ,

operator [])
```

Definition at line 134 of file counters-meat.hpp.

## 9.26.2.13 COUNTERS\_TEMPLATE() [3/9]

## 9.26.2.14 COUNTERS\_TEMPLATE() [4/9]

```
COUNTERS_TEMPLATE (

COUNTERS_TYPE() ,

operator ) const
```

## 9.26.2.15 COUNTERS\_TEMPLATE() [5/9]

```
COUNTERS_TEMPLATE (
          std::vector< double > ,
          gen_hash ) const &
```

#### 9.26.2.16 COUNTERS\_TEMPLATE() [6/9]

```
COUNTERS_TEMPLATE (
          std::vector< std::string > ,
          get_descriptions ) const
```

Definition at line 213 of file counters-meat.hpp.

# 9.26.2.17 COUNTERS\_TEMPLATE() [7/9]

```
COUNTERS_TEMPLATE (
          std::vector< std::string > ,
          get_names ) const
```

Definition at line 201 of file counters-meat.hpp.

# 9.26.2.18 COUNTERS\_TEMPLATE() [8/9]

# 9.26.2.19 **COUNTERS\_TEMPLATE()** [9/9]

# 9.26.2.20 data()

## 9.26.2.21 desc()

Move constructor.

Definition at line 32 of file counters-meat.hpp.

### 9.26.2.22 for()

```
for ( auto &c:data )
```

Definition at line 233 of file counters-meat.hpp.

# 9.26.2.23 hasher() [1/2]

Definition at line 141 of file counters-meat.hpp.

## 9.26.2.24 hasher() [2/2]

Definition at line 144 of file counters-meat.hpp.

## 9.26.2.25 hasher\_fun() [1/2]

Definition at line 13 of file counters-meat.hpp.

#### 9.26.2.26 hasher\_fun() [2/2]

# 9.26.2.27 if() [1/3]

```
if (
    add_dims )
```

Definition at line 248 of file counters-meat.hpp.

## 9.26.2.28 if() [2/3]

```
if ( hasher )
```

Definition at line 255 of file counters-meat.hpp.

```
9.26.2.29 if() [3/3]
```

```
if ( {\tt res.} \quad {\tt size() = =0u \ )}
```

### 9.26.2.30 init\_fun() [1/3]

## 9.26.2.31 init\_fun() [2/3]

## 9.26.2.32 init\_fun() [3/3]

### 9.26.2.33 name()

## 9.26.3 Variable Documentation

#### 9.26.3.1 add\_dims

```
bool add_dims
```

## Initial value:

```
std::vector<double> res
```

Definition at line 227 of file counters-meat.hpp.

#### 9.26.3.2 count\_fun\_

```
Data_Type count_fun_
```

Definition at line 179 of file counters-meat.hpp.

#### 9.26.3.3 counter

```
Data_Type counter

Initial value:
{
    data.push_back(counter)
```

Definition at line 170 of file counters-meat.hpp.

## 9.26.3.4 counter\_

```
Data_Type & counter_

Initial value:
{
    if (this != &counter_) {
        this->count_fun = counter_.count_fun;
        this->init_fun = counter_.init_fun;
        this->hasher_fun = counter_.hasher_fun;

        this->data = counter_.data;
        this->name = counter_.name;
        this->desc = counter_.desc;
    }
    return *this
```

Definition at line 12 of file counters-meat.hpp.

## 9.26.3.5 data\_

Data\_Type Counter\_fun\_type<Array\_Type,Data\_Type> Hasher\_fun\_type<Array\_Type,Data\_Type> Data←
\_Type data\_

Definition at line 182 of file counters-meat.hpp.

#### 9.26.3.6 desc\_

```
Data_Type Counter_fun_type<Array_Type,Data_Type> Hasher_fun_type<Array_Type,Data_Type> Data←
_Type std::string std::string desc_
```

### Initial value:

```
data.emplace_back(Counter<Array_Type,Data_Type>(
    count_fun_,
    init_fun_,
    hasher_fun_,
    data_,
    name_,
    desc_
))
```

Definition at line 184 of file counters-meat.hpp.

#### 9.26.3.7 fun

```
Data_Type fun

Initial value:
{
    hasher_fun = fun
```

Definition at line 111 of file counters-meat.hpp.

#### 9.26.3.8 fun

```
Data_Type fun_
Initial value:
```

hasher = fun\_

Definition at line 270 of file counters-meat.hpp.

## 9.26.3.9 hasher\_fun\_

Data\_Type Counter\_fun\_type<Array\_Type,Data\_Type> Hasher\_fun\_type<Array\_Type,Data\_Type> hasher← \_fun\_

Definition at line 181 of file counters-meat.hpp.

#### 9.26.3.10 i

```
size_t i
```

Definition at line 83 of file counters-meat.hpp.

#### 9.26.3.11 init\_fun\_

```
Data_Type Counter_fun_type<Array_Type,Data_Type> init_fun_
```

Definition at line 180 of file counters-meat.hpp.

#### 9.26.3.12 j

```
size_t j

Initial value:
{
    if (count_fun == nullptr)
        return 0.0
```

Definition at line 83 of file counters-meat.hpp.

#### 9.26.3.13 name\_

```
Data_Type Counter_fun_type<Array_Type,Data_Type> Hasher_fun_type<Array_Type,Data_Type> Data←
_Type std::string name_
```

Definition at line 183 of file counters-meat.hpp.

#### 9.26.3.14 noexcept

```
Data_Type &&counters_ noexcept

Initial value:
{
    if (this != &counter_)
    {
        this->data = std::move(counter_.data);

        this->count_fun = std::move(counter_.count_fun);
        this->init_fun = std::move(counter_.init_fun);
        this->hasher_fun = std::move(counter_.hasher_fun);

        this->name = std::move(counter_.name);
        this->desc = std::move(counter_.desc);
    }
    return *this
```

Definition at line 26 of file counters-meat.hpp.

#### 9.26.3.15 res

return res

Definition at line 265 of file counters-meat.hpp.

#### 9.26.3.16 return

return

Definition at line 175 of file counters-meat.hpp.

# 9.27 include/barry/counters/network-css.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Macros**

- #define CSS\_SIZE()
- #define CSS\_MATCH\_TYPE()
- #define CSS\_CASE\_TRUTH() if ((i < n) && (j < n))
- #define CSS\_TRUE\_CELLS()
- #define CSS\_CASE\_PERCEIVED() else if (((i >= s) && (i < e)) & ((j >= s) && (j < e)))
- #define CSS\_PERCEIVED\_CELLS()
- #define CSS\_CASE\_ELSE()
- #define CSS\_CHECK\_SIZE\_INIT()
- #define CSS\_CHECK\_SIZE()
- #define CSS\_APPEND(name)
- #define CSS\_NET\_COUNTER\_LAMBDA\_INIT()

#### **Functions**

```
• template<typename Tnet = Network>
  void counter css partially false recip commi (NetCounters< Tnet > *counters, size t netsize, const std ←
  ::vector< size_t > &end_, size_t counter_type=0u)
     Counts errors of commission.
• template<typename Tnet = Network>
  void counter css partially false recip omiss (NetCounters< Tnet > *counters, size t netsize, const std↔
  ::vector< size_t > &end_, size_t counter_type=0u)
     Counts errors of omission.
• template<typename Tnet = Network>
  void counter css completely false recip comiss (NetCounters< Tnet > *counters, size t netsize, const
  std::vector< size_t > &end_, size_t counter_type=0u)
     Counts completely false reciprocity (comission)
template<typename Tnet = Network>
  void counter css completely false recip omiss (NetCounters< Tnet > *counters, size t netsize, const
  std::vector< size_t > &end_, size_t counter_type=0u)
     Counts completely false reciprocity (omission)
template<typename Tnet = Network>
  void counter css mixed recip (NetCounters< Tnet > *counters, size t netsize, const std::vector< size t >
  &end , size t counter type=0u)
     Counts mixed reciprocity errors.
• template<typename Tnet = Network>
  void counter_css_census01 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end_, size_t counter_type=0u)

    template<tvpename Tnet = Network>

  void counter_css_census02 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end , size t counter type=0u)

    template<typename Tnet = Network>

  void counter_css_census03 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end_, size_t counter_type=0u)
template<typename Tnet = Network>
  void counter_css_census04 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end_, size_t counter_type=0u)
template<typename Tnet = Network>
  void counter_css_census05 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end_, size_t counter_type=0u)
template<typename Tnet = Network>
  void counter css census06 (NetCounters< Tnet > *counters, size t netsize, const std::vector< size t >
  &end_, size_t counter_type=0u)
template<typename Tnet = Network>
  void counter_css_census07 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end_, size_t counter_type=0u)
template<typename Tnet = Network>
  void counter_css_census08 (NetCounters< Tnet > *counters, size_t netsize, const std::vector< size_t >
  &end , size t counter type=0u)
template<typename Tnet = Network>
  void counter css census09 (NetCounters< Tnet > *counters, size t netsize, const std::vector< size t >
  &end_, size_t counter_type=0u)
template<typename Tnet = Network>
  void counter css census10 (NetCounters< Tnet > *counters, size t netsize, const std::vector< size t >
  &end , size t counter type=0u)
```

#### 9.27.1 Macro Definition Documentation

## 9.27.1.1 CSS\_APPEND

Definition at line 81 of file network-css.hpp.

## 9.27.1.2 CSS\_CASE\_ELSE

```
#define CSS_CASE_ELSE( )
```

Definition at line 66 of file network-css.hpp.

#### 9.27.1.3 CSS CASE PERCEIVED

```
      \# define \ CSS\_CASE\_PERCEIVED( ) \ else \ if \ (((i >= s) \ \&\& \ (i < e)) \ \& \ ((j >= s) \ \&\& \ (j < e)))
```

Definition at line 48 of file network-css.hpp.

## 9.27.1.4 CSS\_CASE\_TRUTH

```
#define CSS_CASE_TRUTH( ) if ((i < n) && (j < n))
```

Definition at line 32 of file network-css.hpp.

#### 9.27.1.5 CSS CHECK SIZE

```
#define CSS_CHECK_SIZE( )

Value:
    for (size_t i = 0u; i < end_.size(); ++i) {\
    if (i == 0u) continue; \
    else if (end_[i] < end_[i-1u]) \</pre>
```

throw std::logic\_error("Endpoints should be specified in order.");}

Definition at line 76 of file network-css.hpp.

## 9.27.1.6 CSS\_CHECK\_SIZE\_INIT

Definition at line 70 of file network-css.hpp.

## 9.27.1.7 CSS\_MATCH\_TYPE

Definition at line 22 of file network-css.hpp.

## 9.27.1.8 CSS\_NET\_COUNTER\_LAMBDA\_INIT

Definition at line 89 of file network-css.hpp.

#### 9.27.1.9 CSS PERCEIVED CELLS

```
#define CSS_PERCEIVED_CELLS()

Value:
    size_t i_ = i - s; \
    size_t j_ = j - s; \
    CSS_MATCH_TYPE() \
    double tji = static_cast<double>(Array(j - s, i - s, false)); \
    double pji = static_cast<double>(Array(j, i, false)); \
    double tij = static_cast<double>(Array(i - s, j - s, false));
```

Definition at line 55 of file network-css.hpp.

## 9.27.1.10 CSS\_SIZE

```
#define CSS_SIZE()

Value:
    size_t n = data.indices[0u]; \
    size_t s = data.indices[1u]; \
    size_t e = data.indices[2u]; \
    size_t ctype = data.indices[3u]; \
    size_t ego_id = data.indices[4u]; \
    if (ctype > 2) \
        throw std::range_error("Counter type should be 0, 1, or 2.");
```

Definition at line 8 of file network-css.hpp.

## 9.27.1.11 CSS\_TRUE\_CELLS

```
#define CSS_TRUE_CELLS()

Value:
    size_t i_ = i; \
    size_t j_ = j; \
    CSS_MATCH_TYPE() \
    double tji = static_cast<double>(Array(j, i, false)); \
    double pij = static_cast<double>(Array(i + s, j + s, false)); \
    double pji = static_cast<double>(Array(j + s, i + s, false));
```

Definition at line 39 of file network-css.hpp.

## 9.27.2 Function Documentation

## 9.27.2.1 counter css census01()

Definition at line 324 of file network-css.hpp.

## 9.27.2.2 counter\_css\_census02()

Definition at line 389 of file network-css.hpp.

## 9.27.2.3 counter\_css\_census03()

Definition at line 429 of file network-css.hpp.

#### 9.27.2.4 counter css census04()

Definition at line 469 of file network-css.hpp.

## 9.27.2.5 counter\_css\_census05()

Definition at line 509 of file network-css.hpp.

## 9.27.2.6 counter\_css\_census06()

Definition at line 549 of file network-css.hpp.

## 9.27.2.7 counter\_css\_census07()

Definition at line 589 of file network-css.hpp.

#### 9.27.2.8 counter css census08()

Definition at line 629 of file network-css.hpp.

## 9.27.2.9 counter\_css\_census09()

Definition at line 669 of file network-css.hpp.

## 9.27.2.10 counter\_css\_census10()

Definition at line 709 of file network-css.hpp.

## 9.27.2.11 counter\_css\_completely\_false\_recip\_comiss()

Counts completely false reciprocity (comission)

Definition at line 200 of file network-css.hpp.

## 9.27.2.12 counter\_css\_completely\_false\_recip\_omiss()

Counts completely false reciprocity (omission)

Definition at line 241 of file network-css.hpp.

#### 9.27.2.13 counter css mixed recip()

Counts mixed reciprocity errors.

Definition at line 282 of file network-css.hpp.

## 9.27.2.14 counter\_css\_partially\_false\_recip\_commi()

Counts errors of commission.

#### **Parameters**

netsize	Size of the reference (true) network
end_	Vector indicating one past the ending index of each network. (see details)
counter_type	Size_t indicating the type of counter to use. Possible values are: 0: Count all, 1: Only count if perceiver is involved, and 2: Only count if perceiver is not involved.

The end\_ parameter should be of length  ${\tt N}$  of networks - 1. It is assumed that the first network ends at netsize.

Definition at line 107 of file network-css.hpp.

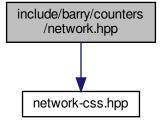
## 9.27.2.15 counter\_css\_partially\_false\_recip\_omiss()

Counts errors of omission.

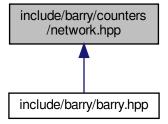
Definition at line 155 of file network-css.hpp.

# 9.28 include/barry/counters/network.hpp File Reference

#include "network-css.hpp"
Include dependency graph for network.hpp:



This graph shows which files directly or indirectly include this file:



#### **Classes**

class NetworkData

Data class for Networks.

· class NetCounterData

Data class used to store arbitrary size\_t or double vectors.

## **Macros**

- #define NET\_C\_DATA\_IDX(i) (data.indices[i])
- #define NET\_C\_DATA\_NUM(i) (data.numbers[i])

## Macros for defining counters

- #define NETWORK\_COUNTER(a)
- #define NETWORK\_COUNTER\_LAMBDA(a)
- #define NETWORKDENSE\_COUNTER\_LAMBDA(a)

## Macros for defining rules

- #define NETWORK\_RULE(a)
- #define NETWORK\_RULE\_LAMBDA(a)

#### **Functions**

```
• template<typename Tnet = Network>
  void counter_edges (NetCounters < Tnet > *counters)
     Number of edges.
template<typename Tnet = Network>
  void counter isolates (NetCounters < Tnet > *counters)
     Number of isolated vertices.

    template<> void counter_isolates (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter mutual (NetCounters < Tnet > *counters)
     Number of mutual ties.
• template<typename Tnet = Network>
  void counter istar2 (NetCounters < Tnet > *counters)

    template<> void counter istar2 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter ostar2 (NetCounters< Tnet > *counters)

    template<> void counter ostar2 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter ttriads (NetCounters< Tnet > *counters)

    template<> void counter_ttriads (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter_ctriads (NetCounters< Tnet > *counters)

    template<> void counter_ctriads (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter_density (NetCounters < Tnet > *counters)
• template<typename Tnet = Network>
  void counter_idegree15 (NetCounters< Tnet > *counters)

    template<> void counter idegree15 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter odegree15 (NetCounters< Tnet > *counters)

    template<> void counter_odegree15 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter_absdiff (NetCounters < Tnet > *counters, size_t attr_id, double alpha=1.0)
     Sum of absolute attribute difference between ego and alter.
• template<typename Tnet = Network>
  void counter diff (NetCounters < Tnet > *counters, size t attr id, double alpha=1.0, double tail head=true)
     Sum of attribute difference between ego and alter to pow(alpha)

    NETWORK_COUNTER (init_single_attr)

    template<typename Tnet = Network>

  void counter nodeicov (NetCounters < Tnet > *counters, size t attr id)
template<typename Tnet = Network>
  void counter nodeocov (NetCounters< Tnet > *counters, size t attr id)
template<typename Tnet = Network>
  void counter_nodecov (NetCounters< Tnet > *counters, size_t attr_id)
template<typename Tnet = Network>
  void counter_nodematch (NetCounters < Tnet > *counters, size_t attr_id)
• template<typename Tnet = Network>
  void counter_idegree (NetCounters< Tnet > *counters, std::vector< size t > d)
     Counts number of vertices with a given in-degree.

    template<> void counter_idegree (NetCounters< NetworkDense > *counters, std::vector< size_t > d)

template<typename Tnet = Network>
  void counter_odegree (NetCounters< Tnet > *counters, std::vector< size_t > d)
     Counts number of vertices with a given out-degree.

    template<> void counter_odegree (NetCounters< NetworkDense > *counters, std::vector< size_t > d)
```

template < typename Tnet = Network>
 void counter\_degree (NetCounters < Tnet > \*counters, std::vector < size\_t > d)
 Counts number of vertices with a given out-degree.

#### Rules for network models

#### **Parameters**

```
rules | A pointer to a NetRules object (Rules < Network, bool > ).
```

template<typename Tnet = Network>
 void rules\_zerodiag (NetRules< Tnet > \*rules)
 Number of edges.

## Convenient typedefs for network objects.

- #define BARRY ZERO NETWORK 0.0
- #define BARRY\_ZERO\_NETWORK\_DENSE 0
- typedef BArray< double, NetworkData > Network
- typedef BArrayDense< int, NetworkData > NetworkDense
- template < typename Tnet = Network > using NetCounter = Counter < Tnet, NetCounterData >
- template < typename Tnet = Network >
   using NetCounters = Counters < Tnet, NetCounterData >
- template<typename Tnet = Network>
   using NetSupport = Support< Tnet, NetCounterData >
- template<typename Tnet = Network>
   using NetStatsCounter = StatsCounter< Tnet, NetCounterData >
- template<typename Tnet >
   using NetModel = Model< Tnet, NetCounterData >
- template<typename Tnet = Network>
   using NetRule = Rule< Tnet, bool >
- template<typename Tnet = Network>
   using NetRules = Rules< Tnet, bool >

#### 9.28.1 Macro Definition Documentation

## 9.28.1.1 BARRY\_ZERO\_NETWORK

#define BARRY ZERO NETWORK 0.0

Definition at line 85 of file network.hpp.

## 9.28.1.2 BARRY\_ZERO\_NETWORK\_DENSE

```
#define BARRY_ZERO_NETWORK_DENSE 0
```

Definition at line 86 of file network.hpp.

## 9.28.1.3 NET\_C\_DATA\_IDX

Definition at line 74 of file network.hpp.

## 9.28.1.4 NET\_C\_DATA\_NUM

Definition at line 75 of file network.hpp.

## 9.28.1.5 NETWORK\_COUNTER

#### Value:

```
template<typename Tnet = Network>\
inline double (a) (const Tnet & Array, size_t i, size_t j, NetCounterData & data)
```

Function for definition of a network counter function

Definition at line 114 of file network.hpp.

## 9.28.1.6 NETWORK\_COUNTER\_LAMBDA

### Value:

```
Counter_fun_type<Tnet, NetCounterData> a = \
   [](const Tnet & Array, size_t i, size_t j, NetCounterData & data)
```

Lambda function for definition of a network counter function

Definition at line 119 of file network.hpp.

## 9.28.1.7 NETWORK\_RULE

#### Value:

```
template<typename Tnet = Network>\
inline bool (a) (const Tnet & Array, size_t i, size_t j, bool & data)
```

Function for definition of a network counter function

Definition at line 133 of file network.hpp.

#### 9.28.1.8 NETWORK RULE LAMBDA

```
#define NETWORK_RULE_LAMBDA(
a )
```

#### Value:

```
Rule_fun_type<Tnet, bool> a = \
[](const Tnet & Array, size_t i, size_t j, bool & data)
```

Lambda function for definition of a network counter function

Definition at line 138 of file network.hpp.

## 9.28.1.9 NETWORKDENSE\_COUNTER\_LAMBDA

```
#define NETWORKDENSE_COUNTER_LAMBDA( a \ )
```

#### Value:

```
Counter_fun_type<NetworkDense, NetCounterData> a = \
   [](const NetworkDense & Array, size_t i, size_t j, NetCounterData & data)
```

Definition at line 123 of file network.hpp.

## 9.28.2 Typedef Documentation

#### 9.28.2.1 NetCounter

```
template<typename Tnet = Network>
using NetCounter = Counter<Tnet, NetCounterData >
```

Definition at line 89 of file network.hpp.

#### 9.28.2.2 NetCounters

```
template<typename Tnet = Network>
using NetCounters = Counters<Tnet, NetCounterData>
```

Definition at line 92 of file network.hpp.

#### 9.28.2.3 NetModel

```
template<typename Tnet >
using NetModel = Model<Tnet, NetCounterData>
```

Definition at line 101 of file network.hpp.

## 9.28.2.4 NetRule

```
template<typename Tnet = Network>
using NetRule = Rule<Tnet, bool>
```

Definition at line 104 of file network.hpp.

#### 9.28.2.5 NetRules

```
template<typename Tnet = Network>
using NetRules = Rules<Tnet, bool>
```

Definition at line 107 of file network.hpp.

#### 9.28.2.6 NetStatsCounter

```
template<typename Tnet = Network>
using NetStatsCounter = StatsCounter<Tnet, NetCounterData>
```

Definition at line 98 of file network.hpp.

## 9.28.2.7 NetSupport

```
template<typename Tnet = Network>
using NetSupport = Support<Tnet, NetCounterData >
```

Definition at line 95 of file network.hpp.

#### 9.28.2.8 Network

```
typedef BArray<double, NetworkData> Network
```

Definition at line 82 of file network.hpp.

### 9.28.2.9 NetworkDense

```
typedef BArrayDense<int, NetworkData> NetworkDense
```

Definition at line 83 of file network.hpp.

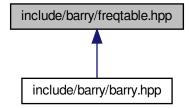
## 9.28.3 Function Documentation

## 9.28.3.1 rules\_zerodiag()

Number of edges.

Definition at line 1381 of file network.hpp.

# 9.29 include/barry/freqtable.hpp File Reference



#### **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:

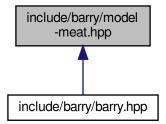


## Classes

- 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



#### **Functions**

- double update\_normalizing\_constant (const std::vector< double > &params, const double \*support, size\_t
   k, size t n)
- double likelihood\_ (const double \*stats\_target, const std::vector< double > &params, const double normalizing\_constant, size\_t n\_params, bool log\_=false)

#### 9.31.1 Function Documentation

## 9.31.1.1 likelihood\_()

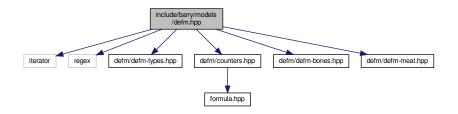
Definition at line 95 of file model-meat.hpp.

## 9.31.1.2 update\_normalizing\_constant()

Definition at line 9 of file model-meat.hpp.

## 9.32 include/barry/models/defm.hpp File Reference

```
#include <iterator>
#include <regex>
#include "defm/defm-types.hpp"
#include "defm/counters.hpp"
#include "defm/defm-bones.hpp"
#include "defm/defm-meat.hpp"
Include dependency graph for defm.hpp:
```

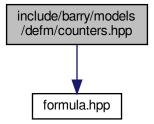


## **Namespaces**

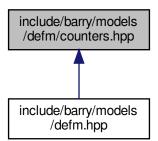
• defm

# 9.33 include/barry/models/defm/counters.hpp File Reference

#include "formula.hpp"
Include dependency graph for counters.hpp:



This graph shows which files directly or indirectly include this file:



## **Macros**

- #define MAKE\_DEFM\_HASHER(hasher, a, cov)
- #define DEFM\_RULEDYN\_LAMBDA(a)
- #define UNI\_SUB(a)

## Macros for defining counters

- #define DEFM\_COUNTER(a) inline double (a) (const DEFMArray & Array, size\_t i, size\_t j, DEFMCounterData & data)
- #define DEFM\_COUNTER\_LAMBDA(a)

#### Macros for defining rules

- #define DEFM\_RULE(a) inline bool (a) (const DEFMArray & Array, size\_t i, size\_t j, bool & data)
- #define DEFM\_RULE\_LAMBDA(a)

#### **Functions**

void counter\_ones (DEFMCounters \*counters, int covar\_index=-1, std::string vname="", const std::vector< std::string > \*x\_names=nullptr)

Prevalence of ones.

- void counter\_logit\_intercept (DEFMCounters \*counters, size\_t n\_y, std::vector< size\_t > which={}, int covar\_index=-1, std::string vname="", const std::vector< std::string > \*x\_names=nullptr, const std::vector< std::string > \*y\_names=nullptr)

Prevalence of ones.

void counter\_transition\_formula (DEFMCounters \*counters, std::string formula, size\_t m\_order, size\_t n\_y, int covar\_index=-1, std::string vname="", const std::vector< std::string > \*x\_names=nullptr, const std::vector< std::string > \*y\_names=nullptr)

Prevalence of ones.

• void counter\_fixed\_effect (DEFMCounters \*counters, int covar\_index, double k, std::string vname="", const std::vector< std::string > \*x names=nullptr)

Prevalence of ones.

### Returns true if the cell is free

### **Parameters**

rules | A pointer to a DEFMRules object (Rules < DEFMArray, bool > ).

- void rules\_markov\_fixed (DEFMRules \*rules, size\_t markov\_order)
  - Number of edges.
- void rules\_dont\_become\_zero (DEFMSupport \*support, std::vector< size\_t > ids)

Blocks switching a one to zero.

## 9.33.1 Macro Definition Documentation

## 9.33.1.1 DEFM\_COUNTER

Function for definition of a network counter function

Definition at line 39 of file counters.hpp.

## 9.33.1.2 DEFM\_COUNTER\_LAMBDA

[](const DEFMArray & Array, size\_t i, size\_t j, DEFMCounterData & data) -> double

Lambda function for definition of a network counter function

Definition at line 43 of file counters.hpp.

## 9.33.1.3 DEFM\_RULE

Function for definition of a network counter function

Definition at line 53 of file counters.hpp.

## 9.33.1.4 DEFM\_RULE\_LAMBDA

#### Value:

```
barry::Rule_fun_type<DEFMArray, DEFMRuleData> a = \
[](const DEFMArray & Array, size_t i, size_t j, DEFMRuleData & data) -> bool
```

Lambda function for definition of a network counter function

Definition at line 57 of file counters.hpp.

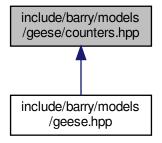
## 9.33.1.5 DEFM\_RULEDYN\_LAMBDA

Lambda function for definition of a network counter function

Definition at line 63 of file counters.hpp.

## 9.33.1.6 UNI\_SUB

# 9.34 include/barry/models/geese/counters.hpp File Reference



#### **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.

- 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.

Cap the number of functions per gene.

- $\bullet \ \ void\ counter\_maxfuns\ (PhyloCounters *counters, size\_t\ lb, size\_t\ ub, size\_t\ duplication = Geese :: etype\_default)$
- 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)
- 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)

Total number of changes. Use this statistic to account for "preservation".

Longest branch mutates (either by gain or by loss)

- void counter\_neofun (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

  Total number of neofunctionalization events.
- void counter\_pairwise\_neofun\_singlefun (PhyloCounters \*counters, size\_t nfunA, size\_t duplication=Geese::etype\_default)

  Total number of neofunctionalization events sum\_u sum\_{{w < u}} [x(u,a)\*(1 x(w,a)) + (1 x(u,a)) \* x(w,a)] change stat: delta{x(u,a): 0-> 1} = 1 2 \* x(w,a)
- void counter\_neofun\_a2b (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

  Total number of neofunctionalization events.

• void counter\_co\_opt (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_t duplication=Geese::etype\_default)

Function co-opting.

- void counter\_k\_genes\_changing (PhyloCounters \*counters, size\_t k, size\_t duplication=Geese::etype\_default)

  Indicator function. Equals to one if k genes changed and zero otherwise.
- void counter\_less\_than\_p\_prop\_genes\_changing (PhyloCounters \*counters, double p, size\_t duplication=Geese::etype\_default Indicator function. Equals to one if k genes changed and zero otherwise.
- void counter\_gains\_from\_0 (PhyloCounters \*counters, std::vector < size\_t > nfun, size\_t duplication=Geese::etype\_default)

  Used when all the functions are in 0 (like the root node prob.)
- void counter\_overall\_gains\_from\_0 (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

  Used when all the functions are in 0 (like the root node prob.)
- void counter\_pairwise\_overall\_change (PhyloCounters \*counters, size\_t duplication=Geese::etype\_default)

  Used when all the functions are in 0 (like the root node prob.)
- void counter\_pairwise\_preserving (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_ t duplication=Geese::etype\_default)

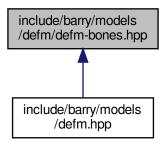
Used when all the functions are in 0 (like the root node prob.)

void counter\_pairwise\_first\_gain (PhyloCounters \*counters, size\_t nfunA, size\_t nfunB, size\_
 t duplication=Geese::etype\_default)

Used when all the functions are in 0 (like the root node prob.)

## 9.35 include/barry/models/defm/defm-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:

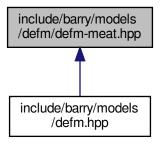


#### **Classes**

• class DEFM

## 9.36 include/barry/models/defm/defm-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



## **Macros**

- #define DEFM\_RANGES(a)
- #define DEFM\_LOOP\_ARRAYS(a) for (size\_t a = 0u; a < (nobs\_i M\_order); ++a)</li>

## **Functions**

• std::vector< double > keygen\_defm (const DEFMArray &Array\_, DEFMCounterData \*data)

## 9.36.1 Macro Definition Documentation

## 9.36.1.1 DEFM\_LOOP\_ARRAYS

Definition at line 36 of file defm-meat.hpp.

## 9.36.1.2 DEFM\_RANGES

Definition at line 31 of file defm-meat.hpp.

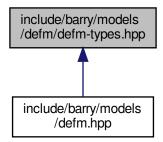
#### 9.36.2 Function Documentation

### 9.36.2.1 keygen\_defm()

Definition at line 4 of file defm-meat.hpp.

## 9.37 include/barry/models/defm/defm-types.hpp File Reference

This graph shows which files directly or indirectly include this file:



### **Classes**

- · class DEFMData
  - Data class for DEFM arrays.
- class DEFMCounterData

Data class used to store arbitrary size\_t or double vectors.

- class DEFMRuleData
- · class DEFMRuleDynData

## **Typedefs**

typedef barry::BArrayDense< int, DEFMData > DEFMArray

## Convenient typedefs for network objects.

- typedef barry::Counter< DEFMArray, DEFMCounterData > DEFMCounter
- typedef barry::Counters < DEFMArray, DEFMCounterData > DEFMCounters
- typedef barry::Support
   DEFMArray, DEFMCounterData, DEFMRuleData, DEFMRuleDynData > DEFMSupport
- typedef barry::StatsCounter< DEFMArray, DEFMCounterData > DEFMStatsCounter
- typedef barry::Model DEFMArray, DEFMCounterData, DEFMRuleData, DEFMRuleDynData >
- $\bullet \ \ type def \ barry:: Rule < DEFMArray, \ DEFMRule Data > DEFMRule \\$
- typedef barry::Rules< DEFMArray, DEFMRuleData > DEFMRules
- typedef barry::Rule < DEFMArray, DEFMRuleDynData > DEFMRuleDyn
- typedef barry::Rules < DEFMArray, DEFMRuleDynData > DEFMRulesDyn

## 9.37.1 Typedef Documentation

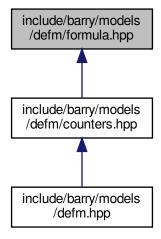
#### 9.37.1.1 **DEFMArray**

typedef barry::BArrayDense<int, DEFMData> DEFMArray

Definition at line 5 of file defm-types.hpp.

# 9.38 include/barry/models/defm/formula.hpp File Reference

This graph shows which files directly or indirectly include this file:



## **Functions**

void defm\_motif\_parser (std::string formula, std::vector< size\_t > &locations, std::vector< bool > &signs, size\_t m\_order, size\_t y\_ncol)

Parses a motif formula.

## 9.38.1 Function Documentation

## 9.38.1.1 defm\_motif\_parser()

Parses a motif formula.

This function will take the formula and generate the corresponding input for defm::counter\_transition(). Formulas can be specified in the following ways:

- Intercept effect: {...} No transition, only including the current state.
- Transition effect: {...} > {...} Includes current and previous states.

The general notation is  $[0]y[column id]_[row id]$ . A preceding 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.

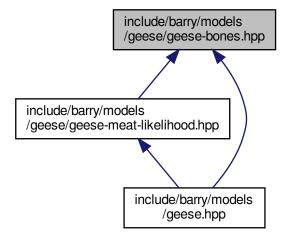
# 9.41 include/barry/models/geese/flock-meat.hpp File Reference

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()

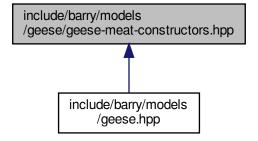
Definition at line 60 of file geese-bones.hpp.

## 9.42.2.4 vector\_caster()

```
template<typename Ta , typename Tb >  $ $ std::vector< Ta > vector\_caster ( \\ const std::vector< Tb > & x ) [inline]
```

Definition at line 10 of file geese-bones.hpp.

# 9.43 include/barry/models/geese/geese-meat-constructors.hpp File Reference



# 9.44 include/barry/models/geese/geese-meat-likelihood.hpp File Reference

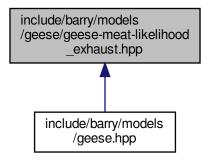
#include "geese-bones.hpp"
Include dependency graph for geese-meat-likelihood.hpp:



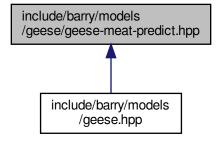


# 9.45 include/barry/models/geese/geese-meat-likelihood\_exhaust.hpp File Reference

This graph shows which files directly or indirectly include this file:

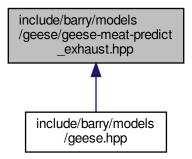


# 9.46 include/barry/models/geese/geese-meat-predict.hpp File Reference



# 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

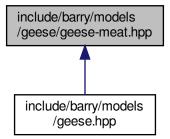


# 9.49 include/barry/models/geese/geese-meat-simulate.hpp File Reference

This graph shows which files directly or indirectly include this file:

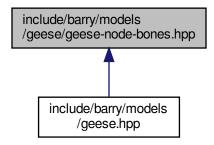


# 9.50 include/barry/models/geese/geese-meat.hpp File Reference



# 9.51 include/barry/models/geese/geese-node-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



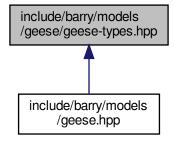
## **Classes**

· 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

#### **Macros**

• #define POS(a, b) (b)\*N + (a)

## **Typedefs**

#### Convenient typedefs for Node objects.

- typedef std::vector< std::pair< size\_t, size\_t >> PhyloRuleData
- typedef barry::BArrayDense< size\_t, NodeData > PhyloArray
- typedef barry::Counter< PhyloArray, PhyloCounterData > PhyloCounter
- typedef barry::Counters < PhyloArray, PhyloCounterData > PhyloCounters
- typedef barry::Rule < PhyloArray, PhyloRuleData > PhyloRule
- typedef barry::Rules
   PhyloArray, PhyloRuleData
   PhyloRules
- typedef barry::Rule
   PhyloArray, PhyloRuleDynData
   PhyloRuleDyn
- typedef barry::Rules< PhyloArray, PhyloRuleDynData > PhyloRulesDyn
- typedef barry::Support
   PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloSupport
- typedef barry::StatsCounter< PhyloArray, PhyloCounterData > PhyloStatsCounter
- typedef barry::Model < PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloModel
- typedef barry::PowerSet < PhyloArray, PhyloRuleData > PhyloPowerSet

## 9.52.1 Macro Definition Documentation

## 9.52.1.1 POS

Definition at line 4 of file geese-types.hpp.

## 9.52.2 Typedef Documentation

#### 9.52.2.1 PhyloArray

```
typedef barry::BArrayDense<size_t, NodeData> PhyloArray
```

Definition at line 104 of file geese-types.hpp.

#### 9.52.2.2 PhyloCounter

typedef barry::Counter<PhyloArray, PhyloCounterData > PhyloCounter

Definition at line 105 of file geese-types.hpp.

#### 9.52.2.3 PhyloCounters

typedef barry::Counters< PhyloArray, PhyloCounterData> PhyloCounters

Definition at line 106 of file geese-types.hpp.

#### 9.52.2.4 PhyloModel

typedef barry::Model<PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloModel

Definition at line 116 of file geese-types.hpp.

## 9.52.2.5 PhyloPowerSet

typedef barry::PowerSet<PhyloArray, PhyloRuleData> PhyloPowerSet

Definition at line 117 of file geese-types.hpp.

#### 9.52.2.6 PhyloRule

typedef barry::Rule<PhyloArray,PhyloRuleData> PhyloRule

Definition at line 108 of file geese-types.hpp.

#### 9.52.2.7 PhyloRuleData

typedef std::vector< std::pair< size\_t, size\_t > > PhyloRuleData

Definition at line 101 of file geese-types.hpp.

#### 9.52.2.8 PhyloRuleDyn

typedef barry::Rule<PhyloArray,PhyloRuleDynData> PhyloRuleDyn

Definition at line 111 of file geese-types.hpp.

#### 9.52.2.9 PhyloRules

typedef barry::Rules<PhyloArray,PhyloRuleData> PhyloRules

Definition at line 109 of file geese-types.hpp.

#### 9.52.2.10 PhyloRulesDyn

typedef barry::Rules<PhyloArray,PhyloRuleDynData> PhyloRulesDyn

Definition at line 112 of file geese-types.hpp.

## 9.52.2.11 PhyloStatsCounter

typedef barry::StatsCounter<PhyloArray, PhyloCounterData> PhyloStatsCounter

Definition at line 115 of file geese-types.hpp.

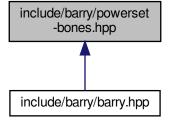
#### 9.52.2.12 PhyloSupport

typedef barry::Support<PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloSupport

Definition at line 114 of file geese-types.hpp.

# 9.53 include/barry/powerset-bones.hpp File Reference

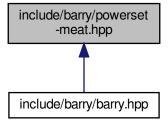
This graph shows which files directly or indirectly include this file:



#### **Classes**

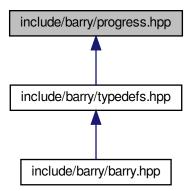
# 9.54 include/barry/powerset-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



# 9.55 include/barry/progress.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Classes**

class Progress

A simple progress bar.

#### **Macros**

• #define BARRY\_PROGRESS\_BAR\_WIDTH 80

#### 9.55.1 Macro Definition Documentation

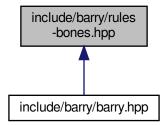
#### 9.55.1.1 BARRY\_PROGRESS\_BAR\_WIDTH

```
#define BARRY_PROGRESS_BAR_WIDTH 80
```

Definition at line 5 of file progress.hpp.

# 9.56 include/barry/rules-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Classes**

class Rule < Array\_Type, Data\_Type >

Rule for determining if a cell should be included in a sequence.

- class Rules < Array\_Type, Data\_Type >

Vector of objects of class Rule.

## **Functions**

template<typename Array\_Type , typename Data\_Type >
 bool rule\_fun\_default (const Array\_Type \*array, size\_t i, size\_t j, Data\_Type \*dat)

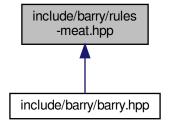
#### 9.56.1 Function Documentation

#### 9.56.1.1 rule\_fun\_default()

Definition at line 5 of file rules-bones.hpp.

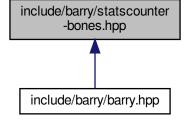
# 9.57 include/barry/rules-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



# 9.58 include/barry/statscounter-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### Classes

 $\bullet \ \ {\it class StatsCounter} < {\it Array\_Type}, \ {\it Data\_Type} >$ 

Count stats for a single Array.

## 9.59 include/barry/statscounter-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Macros**

- #define STATSCOUNTER TYPE() StatsCounter<Array Type, Data Type>
- #define STATSCOUNTER\_TEMPLATE\_ARGS() < typename Array\_Type, typename Data\_Type >
- #define STATSCOUNTER\_TEMPLATE(a, b) template STATSCOUNTER\_TEMPLATE\_ARGS() inline a STATSCOUNTER\_TYPE()::b

## **Functions**

- STATSCOUNTER\_TEMPLATE (, StatsCounter)(const StatsCounter< Array\_Type
- EmptyArray clear ()
- STATSCOUNTER TEMPLATE (,~StatsCounter)()
- STATSCOUNTER\_TEMPLATE (void, reset\_array)(const Array\_Type \*Array\_)
- STATSCOUNTER\_TEMPLATE (void, add\_counter)(Counter< Array\_Type</li>
- STATSCOUNTER\_TEMPLATE (void, set\_counters)(Counters< Array\_Type
- STATSCOUNTER\_TEMPLATE (void, count\_init)(size\_t i
- current\_stats resize (counters->size(), 0.0)
- for (size\_t n=0u;n< counters->size();++n) current\_stats[n]
- STATSCOUNTER\_TEMPLATE (void, count\_current)(size\_t i
- STATSCOUNTER\_TEMPLATE (std::vector< std::string >, get\_names)() const
- STATSCOUNTER\_TEMPLATE (std::vector< std::string >, get\_descriptions)() const

#### **Variables**

```
    Data_Type & counter
    EmptyArray = *Array
    current_stats = counter.current_stats
    counters = new Counters<Array_Type,Data_Type>((*counter.counters))
    counter_deleted = false
    Data_Type f_
    return
```

#### 9.59.1 Macro Definition Documentation

#### 9.59.1.1 STATSCOUNTER\_TEMPLATE

Data\_Type \* counters\_

size\_t j

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:
{
```

throw std::logic\_error("No counters added: Cannot count without knowning what to count!")

Definition at line 66 of file statscounter-meat.hpp.

if (counters->size() == 0u)

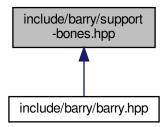
#### 9.59.3.9 return

return

Definition at line 49 of file statscounter-meat.hpp.

# 9.60 include/barry/support-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



#### **Classes**

class Support < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type >
 Compute the support of sufficient statistics.

# 9.61 include/barry/support-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



## **Macros**

• #define BARRY\_SUPPORT\_MEAT\_HPP 1

## 9.61.1 Macro Definition Documentation

## 9.61.1.1 BARRY\_SUPPORT\_MEAT\_HPP

#define BARRY\_SUPPORT\_MEAT\_HPP 1

Definition at line 2 of file support-meat.hpp.

# 9.62 include/barry/typedefs.hpp File Reference

```
#include "barry-configuration.hpp"
#include "barry-debug.hpp"
#include "progress.hpp"
Include dependency graph for typedefs.hpp:
```



This graph shows which files directly or indirectly include this file:



#### **Classes**

- class Entries < Cell\_Type >
  - A wrapper class to store source, target, val from a BArray object.
- struct vecHasher

## **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⇔ _t	Focal cell
Data_Type	Data associated with the function, for example, id of the attribute in the Array.

## Returns

```
Counter_fun_type a double (the change statistic) Rule_fun_type a bool. True if the cell is blocked.
```

Definition at line 187 of file typedefs.hpp.

#### 9.62.1.3 Counts\_type

```
typedef std::vector< std::pair< std::vector<double>, size_t >> Counts_type
Definition at line 51 of file typedefs.hpp.
```

## 9.62.1.4 Hasher\_fun\_type

```
template<typename Array_Type , typename Data_Type >
using Hasher_fun_type = std::function<std::vector<double>(const Array_Type &, Data_Type *)>
```

Hasher function used by the counter.

Used to characterize the support of the array.

#### **Template Parameters**

```
Array_Type
```

Definition at line 200 of file typedefs.hpp.

## 9.62.1.5 MapVec\_type

```
template<typename Ta = double, typename Tb = size_t>
using MapVec_type = std::unordered_map< std::vector< Ta >, Tb, vecHasher<Ta> >
```

Definition at line 128 of file typedefs.hpp.

#### 9.62.1.6 Row\_type

```
template<typename Cell_Type >
using Row_type = Map< size_t, Cell<Cell_Type> >
```

Definition at line 67 of file typedefs.hpp.

## 9.62.1.7 Rule\_fun\_type

```
template<typename Array_Type , typename Data_Type >
using Rule_fun_type = std::function<bool(const Array_Type &, size_t, size_t, Data_Type &)>
```

Definition at line 190 of file typedefs.hpp.

## 9.62.2 Function Documentation

#### 9.62.2.1 sort\_array()

Ascending sorting an array.

It will sort an array solving ties using the next column. Data is stored column-wise.

#### **Template Parameters**



#### **Parameters**



#### Returns

std::vector<size\_t> The sorting index.

Definition at line 141 of file typedefs.hpp.

## 9.62.2.2 vec\_equal()

Compares if -a- and -b- are equal.

#### **Parameters**

a,b Two vectors of the same length

#### Returns

true if all elements are equal.

Definition at line 210 of file typedefs.hpp.

#### 9.62.2.3 vec\_equal\_approx()

Definition at line 235 of file typedefs.hpp.

## 9.62.2.4 vec\_inner\_prod() [1/2]

Definition at line 286 of file typedefs.hpp.

## 9.62.2.5 vec\_inner\_prod() [2/2]

Definition at line 263 of file typedefs.hpp.

# 9.63 README.md File Reference

# Index

```
\simBArray
                                                           NetworkData, 202
    BArray< Cell Type, Data Type >, 63
                                                      \simNode
                                                           Node, 204
\simBArrayCell
    BArrayCell< Cell_Type, Data_Type >, 74
                                                      \simPhyloRuleDynData
~BArrayCell const
                                                           PhyloRuleDynData, 213
    BArrayCell_const< Cell_Type, Data_Type >, 77
                                                      \simPowerSet
{\sim}\mathsf{BArrayDense}
                                                           PowerSet < Array_Type, Data_Rule_Type >, 216
    BArrayDense < Cell_Type, Data_Type >, 83
                                                      \simProgress
\simBArrayDenseCell
                                                           Progress, 221
    BArrayDenseCell< Cell_Type, Data_Type >, 97
                                                      \simRule
\simBArrayRow
                                                           Rule < Array_Type, Data_Type >, 223
    BArrayRow< Cell Type, Data Type >, 110
                                                      \simRules
~BArrayRow const
                                                           Rules < Array Type, Data Type >, 226
    BArrayRow_const < Cell_Type, Data_Type >, 112
                                                      \simStatsCounter
\simBArrayVector
                                                           StatsCounter < Array_Type, Data_Type >, 231
    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 >,
                                                                236
         118
\simCell
                                                      active
    Cell< Cell_Type >, 122
                                                           Cell< Cell_Type >, 125
\simConstBArrayRowIter
    ConstBArrayRowlter< Cell_Type, Data_Type >,
                                                      add
                                                           Cell< Cell_Type >, 123, 124
         127
                                                           FreqTable< T >, 158
\simCounter
                                                      add array
    Counter< Array_Type, Data_Type >, 130
                                                                                       Data Counter_Type,
                                                           Model <
                                                                       Array Type,
\simCounters
                                                                Data_Rule_Type, Data_Rule_Dyn_Type >,
    Counters < Array_Type, Data_Type >, 135
                                                                180
\simDEFMCounterData
                                                      add counter
    Phylo rules, 43
                                                           Counters < Array_Type, Data_Type >, 135, 136
\simDEFMData
                                                           Model<
                                                                       Array_Type,
                                                                                       Data_Counter_Type,
    Phylo rules, 43
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
\simDEFMRuleDynData
                                                                180, 181
    Phylo rules, 43
                                                           StatsCounter< Array_Type, Data_Type >, 231
\simEntries
                                                           Support<
                                                                        Array_Type,
                                                                                       Data_Counter_Type,
    Entries < Cell Type >, 149
                                                                Data_Rule_Type, Data_Rule_Dyn_Type >,
\simFlock
                                                                236
    Flock, 151
                                                      add data
\simFreqTable
                                                           Flock, 151
    FreqTable < T >, 158
                                                      add dims
\simGeese
                                                           counters-meat.hpp, 283
    Geese, 164
                                                      add hash
\simModel
                                                           Counters < Array Type, Data Type >, 136
    Model<
                Array Type,
                                 Data Counter Type,
         Data_Rule_Type, Data_Rule_Dyn_Type >,
                                                      add hasher
                                                           Model<
                                                                                       Data Counter Type,
                                                                       Array Type,
         180
                                                                Data_Rule_Type, Data_Rule_Dyn_Type >,
\simNetCounterData
    NetCounterData, 199
                                                      add_rule
\simNetworkData
```

Model< Array_Type, Data_Counter_Type Data_Rule_Type, Data_Rule_Dyn_Type > 181	
PowerSet< Array_Type, Data_Rule_Type >, 210	6, nrow, 67
217  Pulso < Arroy Type Data Type > 227	operator*=, 67
Rules < Array_Type, Data_Type >, 227 Support < Array_Type, Data_Counter_Type	operator (), 67
	•
Data_Rule_Type, Data_Rule_Dyn_Type >	•
236, 237	operator/=, 69
add_rule_dyn	operator=, 69
Model < Array_Type, Data_Counter_Type	·
Data_Rule_Type, Data_Rule_Dyn_Type >	
181, 182	print, 69
Support< Array_Type, Data_Counter_Type	• —
Data_Rule_Type, Data_Rule_Dyn_Type >	
237	resize, 70
annotations	rm_cell, 70
Node, 205	row, 70
Array	set_data, 71
ConstBArrayRowIter< Cell_Type, Data_Type >	>, swap_cells, 71
127	swap_cols, 71
array	swap_rows, 71
Node, 205	toggle_cell, 72
Phylo rules, 44	toggle_lock, 72
arrays	transpose, 72
Node, 206	visited, 73
arrays2support	zero_col, 72
Model< Array_Type, Data_Counter_Type	
Data_Rule_Type, Data_Rule_Dyn_Type >	
193	BARRAY_TEMPLATE, 248–250
AS_ONE	BARRAY_TEMPLATE_ARGS, 249, 251
EXISTS, 57	BARRAY TYPE, 249, 251
as vector	COL, 249
FreqTable< T >, 158	for, 251
AS ZERO	operator(), 251
EXISTS, 57	rhs, 251
at	ROW, 249
Phylo rules, 39	this, 252
PhyloCounterData, 210	barray-meat.hpp
Thylocounterbata, 210	COL, 252
BArray	ROW, 253
BArray< Cell_Type, Data_Type >, 62, 63	BARRAY_TEMPLATE
BArray< Cell_Type, Data_Type >, 59	
$\sim$ BArray, 63	barray-meat-operators.hpp, 248–250 BARRAY TEMPLATE ARGS
BArray, 62, 63	
BArrayCell< Cell_Type, Data_Type >, 73	barray-meat-operators.hpp, 249, 251
BArrayCell const< Cell Type, Data Type >, 73	BARRAY_TYPE
clear, 63	barray-meat-operators.hpp, 249, 251
col, 63	BArrayCell
D, 64	BArrayCell< Cell_Type, Data_Type >, 74
	BArrayCell< Cell_Type, Data_Type >, 74
D_ptr, 64	~BArrayCell, 74
default_val, 64	BArray< Cell_Type, Data_Type >, 73
flush_data, 64	BArrayCell, 74
get_cell, 64	operator Cell_Type, 75
get_col_vec, 65	operator*=, 75
get_entries, 65	operator+=, 75
get_row_vec, 65	operator-=, 75
insert_cell, 66	operator/=, 75
is_dense, 66	

75	II 00
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
$\sim$ 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, 255, 257
BArrayDense	BDENSE_TEMPLATE_ARGS, 255, 257
BArrayDense < Cell_Type, Data_Type >, 82, 83	BDENSE_TYPE, 256, 258
BArrayDense< Cell_Type, Data_Type >, 79	COL, 256
$\sim$ BArrayDense, 83	POS, 256
BArrayDense, 82, 83	POS_N, 256
BArrayDenseCell< Cell_Type, Data_Type >, 95,	ROW, 256
99	barraydense-meat.hpp
BArrayDenseCol< Cell_Type, Data_Type >, 95,	COL, 258
102	POS, 258
BArrayDenseCol_const< Cell_Type, Data_Type >,	POS_N, 259
95	ROW, 259
BArrayDenseRow< Cell_Type, Data_Type >, 95,	ZERO_CELL, 259
106	BArrayDenseCell
BArrayDenseRow_const< Cell_Type, Data_Type	BArrayDenseCell< Cell_Type, Data_Type >, 97
	BArrayDenseCell< Cell_Type, Data_Type >, 96
>, 95	
clear, 83	~BArrayDenseCell, 97
col, 84	BArrayDense < Cell_Type, Data_Type >, 95, 99
colsum, 84	BArrayDenseCell, 97
D, 84	BArrayDenseCol< Cell_Type, Data_Type >, 99,
D_ptr, 85	102
default_val, 85	BArrayDenseCol_const< Cell_Type, Data_Type >,
get_cell, 85	99, 104
get_col_vec, 85, 86	BArrayDenseRow< Cell_Type, Data_Type >, 106
get_data, 86	BArrayDenseRow_const< Cell_Type, Data_Type
get_entries, 86	>, 109
get_row_vec, 86, 87	operator Cell_Type, 97
insert_cell, 87	operator*=, 97
is_dense, 87	operator+=, 98
is empty, 88	operator-=, 98
ncol, 88	operator/=, 98
nnozero, 88	operator=, 98
	·
nrow, 88	operator==, 99
operator*=, 89	barraydensecell-bones.hpp
operator(), 88, 89	POS, 260
operator+=, 89	barraydensecell-meat.hpp
operator-=, 90	POS, 260
operator/=, 90	BArrayDenseCell_const< Cell_Type, Data_Type >, 100
operator=, 90, 91	BArrayDenseCol< Cell_Type, Data_Type >, 102
operator==, 91	BArrayDenseCol_const< Cell_Type, Data_Type >,
out_of_range, 91	104
print, 91	BArrayDenseRow< Cell_Type, Data_Type >, 106
reserve, 91	BArrayDenseRow_const< Cell_Type, Data_Type
resize, 92	>, 109
•	•

BArrayDenseCol BArrayDenseCol< Cell_Type, Data_Type >, 100 BArrayDenseCol< Cell_Type, Data_Type >, 100	end, 108 operator(), 108 size, 108
BArrayDense < Cell_Type, Data_Type >, 95, 102	BArrayRow
BArrayDenseCell< Cell_Type, Data_Type >, 99, 102	BArrayRow< Cell_Type, Data_Type >, 110 BArrayRow< Cell Type, Data Type >, 109
BArrayDenseCell_const< Cell_Type, Data_Type	~BArrayRow, 110
>, 102	BArrayRow, 110
BArrayDenseCol, 100	operator BArrayRow< Cell_Type, Data_Type >,
begin, 101	110
end, 101 operator(), 101	operator*=, 110 operator+=, 110
size, 101	operator-=, 110
barraydensecol-bones.hpp	operator/=, 111
POS, 261	operator=, 111
POS_N, 261	operator==, 111
ZERO_CELL, 262	barrayrow-meat.hpp
BArrayDenseCol_const	BROW_TEMPLATE, 264, 265 BROW_TEMPLATE_ARGS, 264
BArrayDenseCol_const< Cell_Type, Data_Type >, 103	BROW_TYPE, 264
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 BArrayDenseCol_const, 103	operator BArrayRow_const< Cell_Type, Data_Type >, 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 BArrayDense< Cell_Type, Data_Type >, 95, 106	BArrayVector
BArrayDenseCell< Cell_Type, Data_Type >, 30, 100	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 is_row, 115
operator(), 106 size, 106	operator std::vector< Cell_Type >, 116
barraydenserow-bones.hpp	operator*=, 116
POS, 263	operator+=, 116
POS_N, 263	operator-=, 116
ZERO_CELL, 263	operator/=, 116
BArrayDenseRow_const	operator=, 117
BArrayDenseRow_const< Cell_Type, Data_Type >, 107	operator==, 117 size, 117
BArrayDenseRow_const< Cell_Type, Data_Type >,	BArrayVector_const
107	BArrayVector_const< Cell_Type, Data_Type >,
BArrayDense < Cell_Type, Data_Type >, 95	118
${\sf BArrayDenseCell} {< Cell\_Type, Data\_Type >}, {\color{red} 109}$	BArrayVector_const< Cell_Type, Data_Type >, 117
BArrayDenseCell_const< Cell_Type, Data_Type	~BArrayVector_const, 118
>, 109 BArrayDenseRow_const, 107	BArrayVector_const, 118 begin, 119
begin, 108	end, 119
<b>∵</b>	•

is_col, 119	BARRY_SUPPORT_MEAT_HPP
is_row, 119	support-meat.hpp, 336
operator std::vector< Cell_Type >, 119	BARRY_UNUSED
operator!=, 119	barry-macros.hpp, 270
operator<, 120	BARRY_VERSION
operator<=, 120	barry.hpp, 272
operator>, 120	BARRY_VERSION_MAYOR
operator>=, 120	barry.hpp, 272
operator==, 120	BARRY_VERSION_MINOR
size, 121	barry.hpp, 272
barry, 55	BARRY_ZERO
barry-configuration.hpp	barry-macros.hpp, 270
BARRY_CHECK_SUPPORT, 267	BARRY_ZERO_DENSE
BARRY_ISFINITE, 267	barry-macros.hpp, 270
BARRY_MAX_NUM_ELEMENTS, 267	BARRY_ZERO_NETWORK
BARRY_SAFE_EXP, 267	network.hpp, 298
Map, 268	BARRY_ZERO_NETWORK_DENSE
printf_barry, 267	network.hpp, 298
barry-debug.hpp	BDENSE_TEMPLATE
BARRY DEBUG LEVEL, 268	barraydense-meat-operators.hpp, 255, 257
barry-macros.hpp	BDENSE_TEMPLATE_ARGS
BARRY_NCORES_ARG, 269	barraydense-meat-operators.hpp, 255, 257
BARRY_ONE, 269	BDENSE TYPE
BARRY_ONE_DENSE, 269	barraydense-meat-operators.hpp, 256, 258
BARRY_UNUSED, 270	begin
BARRY_ZERO, 270	BArrayDenseCol< Cell_Type, Data_Type >, 101
BARRY_ZERO_DENSE, 270	BArrayDenseCol_const< Cell_Type, Data_Type >,
barry.hpp	103
BARRY_HPP, 272	BArrayDenseRow< Cell_Type, Data_Type >, 105
BARRY_VERSION, 272	BArrayDenseRow_const< Cell_Type, Data_Type
BARRY_VERSION_MAYOR, 272	>, 108
BARRY_VERSION_MINOR, 272	BArrayVector< Cell_Type, Data_Type >, 115
COUNTER_FUNCTION, 272	BArrayVector_const< Cell_Type, Data_Type >,
COUNTER_LAMBDA, 272	119
RULE_FUNCTION, 273	PhyloCounterData, 210
RULE_LAMBDA, 273	PowerSet < Array_Type, Data_Rule_Type >, 217
barry::counters, 55	Rules < Array_Type, Data_Type >, 227
barry::counters; 55 barry::counters::network, 56	blengths
BARRY CHECK SUPPORT	NodeData, 209
barry-configuration.hpp, 267	BOTH
BARRY_DEBUG_LEVEL	CHECK, 56
barry-debug.hpp, 268	EXISTS, 57
BARRY_HPP	BROW_TEMPLATE
barry.hpp, 272	barrayrow-meat.hpp, 264, 265
BARRY_ISFINITE	BROW_TEMPLATE_ARGS
barry-configuration.hpp, 267	barrayrow-meat.hpp, 264
BARRY_MAX_NUM_ELEMENTS	BROW_TYPE
barry-configuration.hpp, 267	barrayrow-meat.hpp, 264
BARRY_NCORES_ARG	calc
barry-macros.hpp, 269	PowerSet < Array_Type, Data_Rule_Type >, 217
BARRY_ONE	Support< Array_Type, Data_rule_Type >, 217
barry-macros.hpp, 269	Data_Rule_Type, Data_Rule_Dyn_Type >,
BARRY_ONE_DENSE	237
barry-macros.hpp, 269	
BARRY_PROGRESS_BAR_WIDTH	calc_reduced_sequence
progress.hpp, 328	Geese, 164
BARRY_SAFE_EXP	calc_sequence
barry-configuration.hpp, 267	Geese, 165
	Cell

Cell< Cell_Type >, 122, 123 Cell< Cell_Type >, 121	iter, 128 coordiantes_n_free
~Cell, 122 active, 125	Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,
add, 123, 124	241
Cell, 122, 123 operator Cell_Type, 124	<pre>coordiantes_n_locked     Support&lt; Array_Type, Data_Counter_Type,</pre>
operator!=, 124	Data_Rule_Type, Data_Rule_Dyn_Type >,
operator=, 124, 125	242
operator==, 125	coordinates_free
value, 125	PowerSet< Array_Type, Data_Rule_Type >, 219
visited, 125	Support< Array_Type, Data_Counter_Type,
Cell_const< Cell_Type >, 126	Data_Rule_Type, Data_Rule_Dyn_Type >,
change_stats	242
Support< Array_Type, Data_Counter_Type,	coordinates_locked
Data_Rule_Type, Data_Rule_Dyn_Type >,	PowerSet < Array_Type, Data_Rule_Type >, 219
241	Support< Array_Type, Data_Counter_Type,
CHECK, 56	Data_Rule_Type, Data_Rule_Dyn_Type >,
BOTH, 56	242
NONE, 56	Count
ONE, 56	Counter< Array_Type, Data_Type >, 131
TWO, 56 clear	count_all StatsCounter< Array_Type, Data_Type >, 231
BArray< Cell_Type, Data_Type >, 63	count_current
BArrayDense < Cell_Type, Data_Type >, 83	StatsCounter< Array_Type, Data_Type >, 232
FreqTable < T >, 158	count_fun
statscounter-meat.hpp, 331	Counter< Array_Type, Data_Type >, 132
COL	counters-meat.hpp, 278
barray-meat-operators.hpp, 249	count_fun_
barray-meat.hpp, 252	counters-meat.hpp, 283
barraydense-meat-operators.hpp, 256	count_init
barraydense-meat.hpp, 258	StatsCounter< Array_Type, Data_Type >, 232
col	Counter
BArray < Cell_Type, Data_Type >, 63	Counter< Array_Type, Data_Type >, 130
BArrayDense < Cell_Type, Data_Type >, 84	counter
Col_type	counters-meat.hpp, 284
typedefs.hpp, 339 colnames	statscounter-meat.hpp, 333 Counter< Array_Type, Data_Type >, 128
Flock, 152	~Counter, 130
Geese, 165	count, 131
Model< Array_Type, Data_Counter_Type,	count_fun, 132
Data_Rule_Type, Data_Rule_Dyn_Type >,	Counter, 130
182	data, 132
colsum	desc, 133
BArrayDense < Cell_Type, Data_Type >, 84	get_description, 131
conditional_prob	get_hasher, 131
Model< Array_Type, Data_Counter_Type,	get_name, 131
Data_Rule_Type, Data_Rule_Dyn_Type >,	hasher_fun, 133
182	init, 131
ConstBArrayRowlter	init_fun, 133
ConstBArrayRowlter< Cell_Type, Data_Type >, 127	name, 133
ConstBArrayRowIter< Cell_Type, Data_Type >, 126	operator=, 131, 132 set_hasher, 132
~ConstBArrayRowlter, 127	counter_
Array, 127	counters-meat.hpp, 284
ConstBArrayRowlter, 127	counter_absdiff
current_col, 128	Network counters, 26
current_row, 128	counter_co_opt

Operation 47	Occupio v. 40
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, 291	Counting, 19
counter_css_census02	Phylo counters, 49
network-css.hpp, 291	counter_genes_changing
counter_css_census03	Counting, 19
network-css.hpp, 291	Phylo counters, 50 counter idegree
counter_css_census04 network-css.hpp, 292	Network counters, 29
• •	
counter_css_census05 network-css.hpp, 292	counter_idegree15  Network counters, 29
• •	
counter_css_census06 network-css.hpp, 292	counter_isolates  Network counters, 29, 30
• •	• •
counter_css_census07 network-css.hpp, 292	counter_istar2  Network counters, 30
	counter k genes changing
counter_css_census08 network-css.hpp, 293	Counting, 19
counter css census09	Phylo counters, 50
network-css.hpp, 293	COUNTER_LAMBDA
counter_css_census10	barry.hpp, 272
network-css.hpp, 293	counter_less_than_p_prop_genes_changing
counter_css_completely_false_recip_comiss	Counting, 19
network-css.hpp, 293	Phylo counters, 50
counter_css_completely_false_recip_omiss	counter_logit_intercept
network-css.hpp, 294	Network counters, 30
counter_css_mixed_recip	counter_longest
network-css.hpp, 294	Counting, 20
counter_css_partially_false_recip_commi	Phylo counters, 50
network-css.hpp, 294	counter_loss
counter_css_partially_false_recip_omiss	Counting, 20
network-css.hpp, 295	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, 334	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
193	counter_nodeocov
Counter_fun_type	Network counters, 31
typedefs.hpp, 339	counter_odegree
COUNTER_FUNCTION	Network counters, 31, 32
barry.hpp, 272	counter_odegree15
counter_gains	Network counters, 32

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, 283
Counting, 21	count fun, 278
Phylo counters, 52	count_fun_, 283
counter_overall_loss	counter, 284
Counting, 22	counter , 284
Phylo counters, 52	COUNTER TEMPLATE, 276, 278, 279
counter pairwise first gain	COUNTER TEMPLATE ARGS, 276
Counting, 22	COUNTER TYPE, 277
Phylo counters, 53	COUNTERS TEMPLATE, 277, 279–281
counter pairwise neofun singlefun	COUNTERS TEMPLATE ARGS, 277
Counting, 22	COUNTERS TYPE, 277
Phylo counters, 53	data, 281
counter_pairwise_overall_change	data_, 284
Counting, 22	desc, 281
Phylo counters, 53	desc_, 284
counter_pairwise_preserving	for, 281
Counting, 23	fun, 285
Phylo counters, 53	fun_, 285
counter_preserve_pseudogene	hasher, 281, 282
Counting, 23	hasher_fun, 282
Phylo counters, 54	hasher_fun_, 285
counter_prop_genes_changing	i, 285
Counting, 23	if, 282
Phylo counters, 54	init_fun, 283
counter_subfun	init_fun_, 286
Counting, 23	j, 286
Phylo counters, 54	name, 283
COUNTER_TEMPLATE	name_, 286
counters-meat.hpp, 276, 278, 279	noexcept, 286
COUNTER_TEMPLATE_ARGS	res, 286
counters-meat.hpp, 276	return, 287
counter_transition	TMP_HASHER_CALL, 277
Network counters, 33	counters.hpp
counter_transition_formula	DEFM_COUNTER, 306
Network counters, 34	DEFM_COUNTER_LAMBDA, 307
counter_ttriads	DEFM_RULE, 307
Network counters, 34	DEFM_RULE_LAMBDA, 307
COUNTER_TYPE	DEFM_RULEDYN_LAMBDA, 307
counters-meat.hpp, 277	UNI SUB, 308
Counters	counters_
Counters< Array_Type, Data_Type >, 134, 135	statscounter-meat.hpp, 334
counters	COUNTERS_TEMPLATE
Model< Array_Type, Data_Counter_Type,	counters-meat.hpp, 277, 279–281
Data_Rule_Type, Data_Rule_Dyn_Type >,	COUNTERS_TEMPLATE_ARGS
193	counters-meat.hpp, 277
statscounter-meat.hpp, 334	COUNTERS_TYPE
Counters< Array_Type, Data_Type >, 134	counters-meat.hpp, 277
~Counters. 135	Counting, 13

counter_co_opt, 17	CSS_MATCH_TYPE
counter_cogain, 18	network-css.hpp, 290
counter gains, 18	CSS NET COUNTER LAMBDA INIT
counter_gains_from_0, 18	network-css.hpp, 290
counter_gains_k_offspring, 19	CSS_PERCEIVED_CELLS
counter_genes_changing, 19	network-css.hpp, 290
counter_k_genes_changing, 19	CSS SIZE
counter_less_than_p_prop_genes_changing, 19	network-css.hpp, 290
counter longest, 20	CSS_TRUE_CELLS
counter_loss, 20	network-css.hpp, 291
counter_maxfuns, 20	current_col
counter_neofun, 20	ConstBArrayRowIter< Cell_Type, Data_Type >,
counter_neofun_a2b, 21	128
counter_overall_changes, 21	current_row
counter_overall_gains, 21	ConstBArrayRowIter< Cell_Type, Data_Type >,
counter_overall_gains_from_0, 21	128
counter_overall_loss, 22	current_stats
counter_pairwise_first_gain, 22	statscounter-meat.hpp, 334
counter_pairwise_neofun_singlefun, 22	Support< Array_Type, Data_Counter_Type,
counter_pairwise_overall_change, 22	Data_Rule_Type, Data_Rule_Dyn_Type >,
counter_pairwise_preserving, 23	242
counter_preserve_pseudogene, 23	_
counter_prop_genes_changing, 23	D
counter_subfun, 23	BArray< Cell_Type, Data_Type >, 64
get_last_name, 24	BArrayDense < Cell_Type, Data_Type >, 84
IF_MATCHES, 15	Rule < Array_Type, Data_Type >, 224
IF_NOTMATCHES, 15	D_ptr
IS_DUPLICATION, 15	BArray < Cell_Type, Data_Type >, 64
IS_EITHER, 15	BArrayDense< Cell_Type, Data_Type >, 85
IS_SPECIATION, 16	dat
MAKE_DEFM_HASHER, 16	Flock, 156
MAKE_DUPL_VARS, 16	data
PHYLO_CHECK_MISSING, 16	Counter< Array_Type, Data_Type >, 132
PHYLO_COUNTER_LAMBDA, 17	counters-meat.hpp, 281
PHYLO_RULE_DYN_LAMBDA, 17	PowerSet < Array_Type, Data_Rule_Type >, 219
counts	data_
Phylo rules, 44	counters-meat.hpp, 284
PhyloRuleDynData, 213	default_val BArray< Cell_Type, Data_Type >, 64
Counts_type	BArrayDense < Cell_Type, Data_Type >, 85
typedefs.hpp, 339	DEFM, 139
covar_sort	DEFM, 140
Phylo rules, 44	get_column_major, 140
covar_used	get ID, 140
Phylo rules, 44	get_m_order, 140
covariates	get_model, 141
Phylo rules, 44	get n covars, 141
CSS_APPEND	get_n_obs, 141
network-css.hpp, 288	get_n_rows, 141
CSS_CASE_ELSE	get_n_y, 141
network-css.hpp, 289	get_X, 141
CSS_CASE_PERCEIVED	get_X_names, 142
network-css.hpp, 289	get_Y, 142
CSS_CASE_TRUTH	get_Y_names, 142
network-css.hpp, 289	init, 142
CSS_CHECK_SIZE	is_motif, 142
network-css.hpp, 289	logodds, 142
CSS_CHECK_SIZE_INIT	motif_census, 143
network-css.hpp, 289	print, 143
	1 7 -

set_names, 143	Data_Rule_Type, Data_Rule_Dyn_Type >,
simulate, 143	242
defm, 57	delete_rengine
defm-meat.hpp	Geese, 172
DEFM_LOOP_ARRAYS, 311	Model< Array_Type, Data_Counter_Type,
DEFM_RANGES, 311	Data_Rule_Type, Data_Rule_Dyn_Type >,
keygen_defm, 312	193
defm-types.hpp	delete_rules
DEFMArray, 313	Model < Array_Type, Data_Counter_Type,
DEFM_COUNTER	Data_Rule_Type, Data_Rule_Dyn_Type >,
counters.hpp, 306	194
DEFM_COUNTER_LAMBDA	Support< Array_Type, Data_Counter_Type,
counters.hpp, 307	Data_Rule_Type, Data_Rule_Dyn_Type >,
DEFM_LOOP_ARRAYS	243
defm-meat.hpp, 311	delete_rules_dyn
defm_motif_parser	Model < Array_Type, Data_Counter_Type,
formula.hpp, 313	Data_Rule_Type, Data_Rule_Dyn_Type >,
DEFM RANGES	194
defm-meat.hpp, 311	Support< Array_Type, Data_Counter_Type,
DEFM RULE	Data Rule Type, Data Rule Dyn Type >,
counters.hpp, 307	243
DEFM_RULE_LAMBDA	delete_support
counters.hpp, 307	Geese, 172
DEFM RULEDYN LAMBDA	desc
counters.hpp, 307	Counter< Array_Type, Data_Type >, 133
DEFMArray	counters-meat.hpp, 281
defm-types.hpp, 313	desc_
DEFMCounter	counters-meat.hpp, 284
Phylo rules, 37	directed
DEFMCounterData, 144	NetworkData, 202
Phylo rules, 39	duplication
DEFMCounters	Node, 206
Phylo rules, 37	Node, 200 NodeData, 209
DEFMData, 144	•
,	PhyloRuleDynData, 213
Phylo rules, 39 DEFMModel	empty
	PhyloCounterData, 210
Phylo rules, 37 DEFMRule	EmptyArray
	PowerSet < Array_Type, Data_Rule_Type >, 219
Phylo rules, 38	statscounter-meat.hpp, 334
DEFMRuleData, 146	end
Phylo rules, 40	BArrayDenseCol< Cell_Type, Data_Type >, 101
DEFMRuleDyn  Phylography 200	BArrayDenseCol_const< Cell_Type, Data_Type >,
Phylo rules, 38	103
DEFMRuleDynData, 147	BArrayDenseRow< Cell_Type, Data_Type >, 105
Phylo rules, 40	BArrayDenseRow const< Cell Type, Data Type
DEFMRules	>, 108
Phylo rules, 38	BArrayVector< Cell_Type, Data_Type >, 115
DEFMRulesDyn	BArrayVector_const< Cell_Type, Data_Type >,
Phylo rules, 38	119
DEFMStatsCounter	
Phylo rules, 38	PhyloCounterData, 211
DEFMSupport	PowerSet < Array_Type, Data_Rule_Type >, 217
Phylo rules, 38	Progress, 222
delete_counters	Rules < Array_Type, Data_Type >, 227
Model< Array_Type, Data_Counter_Type,	Entries Call Type > 140
Data_Rule_Type, Data_Rule_Dyn_Type >,	Entries < Cell_Type >, 148
193	Entries < Cell_Type >, 148
Support< Array_Type, Data_Counter_Type,	~Entries, 149
	Entrice 1/18

resize, 149	BArray< Cell_Type, Data_Type >, 64
source, 149	for
target, 149	barray-meat-operators.hpp, 251
val, 149	counters-meat.hpp, 281
etype_default	statscounter-meat.hpp, 331
Geese, 172	formula.hpp
etype_duplication	defm_motif_parser, 313
Geese, 172	FreqTable
etype_either	FreqTable< T >, 158
Geese, 172	FreqTable < T >, 157
etype_speciation	∼FreqTable, 158
Geese, 173	add, 158
eval_rules_dyn	as_vector, 158
Support< Array_Type, Data_Counter_Type,	clear, 158
Data_Rule_Type, Data_Rule_Dyn_Type >,	FreqTable, 158
238	get_data, 159
EXISTS, 57	get_index, 159
AS ONE, 57	make_hash, 159
AS ZERO, 57	print, 159
BOTH, 57	reserve, 159
NONE, 58	size, 160
ONE, 58	fun
TWO, 58	counters-meat.hpp, 285
UKNOWN, 58	fun_
OKNOWN, 30	
f_	counters-meat.hpp, 285
statscounter-meat.hpp, 335	Geese, 160
first_calc_done	∼Geese, 164
Model< Array_Type, Data_Counter_Type,	calc_reduced_sequence, 164
Data_Rule_Type, Data_Rule_Dyn_Type >,	calc_sequence, 165
194	colnames, 165
Flock, 150	delete_rengine, 172
~Flock, 151	delete_support, 172
add_data, 151	etype_default, 172
colnames, 152	etype_duplication, 172
dat, 156	etype_dither, 172
Flock, 151	etype_speciation, 173
get_counters, 152	Geese, 163, 164
get_model, 152	get_annotated_nodes, 165
get_stats_support, 152	get annotations, 165
get_stats_target, 152	get_counters, 165
	get_model, 166
get_support_fun, 153	get probabilities, 166
init, 153	• —
initialized, 156	get_rengine, 166
likelihood_joint, 153	get_states, 166
model, 156	get_support_fun, 166
nfunctions, 156	inherit_support, 167
nfuns, 153	init, 167
nleafs, 154	init_node, 167
nnodes, 154	initialized, 173
nterms, 154	likelihood, 167
ntrees, 154	likelihood_exhaust, 167
operator(), 154	map_to_state_id, 173
parse_polytomies, 155	nannotations, 168
print, 155	nfunctions, 173
rengine, 156	nfuns, 168
set_seed, 155	nleafs, 168
support_size, 155	nnodes, 168
flush_data	nodes, 173

nterms, 168	BArray< Cell_Type, Data_Type >, 65
observed_counts, 169	BArrayDense < Cell_Type, Data_Type >, 85, 86
operator=, 169	get_column_major
parse_polytomies, 169	DEFM, 140
predict, 169	get_counters
predict_backend, 170	Flock, 152
predict_exhaust, 170	Geese, 165
predict_exhaust_backend, 170	Model < Array_Type, Data_Counter_Type,
predict_sim, 170	Data_Rule_Type, Data_Rule_Dyn_Type >,
print, 170	183
print_nodes, 171	PhyloCounterData, 211
print_observed_counts, 171	StatsCounter< Array_Type, Data_Type >, 232
pset_loc, 173	Support< Array_Type, Data_Counter_Type,
reduced_sequence, 174	Data_Rule_Type, Data_Rule_Dyn_Type >,
sequence, 174	238
set_seed, 171	get_counts
simulate, 171	Support< Array_Type, Data_Counter_Type,
support_size, 171	Data Rule Type, Data Rule Dyn Type >,
update_annotations, 171	238
geese, 58	get_current_stats
geese-bones.hpp	Support< Array_Type, Data_Counter_Type,
INITIALIZED, 317	Data_Rule_Type, Data_Rule_Dyn_Type >,
keygen_full, 317	238
RULE_FUNCTION, 317	get_data
vec_diff, 317	BArrayDense < Cell_Type, Data_Type >, 86
vector_caster, 318	FreqTable < T >, 159
geese-types.hpp	PowerSet< Array_Type, Data_Rule_Type >, 217
PhyloArray, 324	Support< Array_Type, Data_Counter_Type,
PhyloCounter, 324	Data_Rule_Type, Data_Rule_Dyn_Type >,
PhyloCounters, 325	239
PhyloModel, 325	get_data_ptr
PhyloPowerSet, 325	PowerSet < Array_Type, Data_Rule_Type >, 218
PhyloRule, 325	get_description
PhyloRuleData, 325	Counter< Array_Type, Data_Type >, 131
PhyloRuleDyn, 325	Rule < Array_Type, Data_Type >, 224
PhyloRules, 326	get_descriptions
PhyloRulesDyn, 326	Counters< Array_Type, Data_Type >, 136
PhyloStatsCounter, 326	Rules< Array_Type, Data_Type >, 227
PhyloSupport, 326	StatsCounter< Array_Type, Data_Type >, 232
POS, 324	get_entries
gen_hash	BArray< Cell_Type, Data_Type >, 65
Counters < Array_Type, Data_Type >, 136	BArrayDense < Cell_Type, Data_Type >, 86
gen_key	get_hasher
Model < Array_Type, Data_Counter_Type,	Counter< Array_Type, Data_Type >, 131
Data_Rule_Type, Data_Rule_Dyn_Type >,	get_ID
183	DEFM, 140
get_annotated_nodes	get_index
Geese, 165	FreqTable $<$ T $>$ , 159
get_annotations	get_last_name
Geese, 165	Counting, 24
get_arrays2support	get_m_order
Model< Array_Type, Data_Counter_Type,	DEFM, 140
Data_Rule_Type, Data_Rule_Dyn_Type >,	get_model
183	DEFM, 141
get cell	Flock, 152
BArray< Cell_Type, Data_Type >, 64	Geese, 166
BArrayDense < Cell_Type, Data_Type >, 85	
get_col_vec	get_n_covars DEFM, 141
901 001 100	DEI IVI, ITI

get n obs	239
DEFM, 141	get seq
get_n_rows	Rules < Array_Type, Data_Type >, 228
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 >, 224	Data_Rule_Type, Data_Rule_Dyn_Type >
get_names	186
Counters< Array_Type, Data_Type >, 137	get_stats_target
Rules < Array_Type, Data_Type >, 228	Flock, 152
StatsCounter< Array_Type, Data_Type >, 232	Model < Array_Type, Data_Counter_Type,
get_norm_const  Model< Array Type, Data Counter Type,	Data_Rule_Type, Data_Rule_Dyn_Type >: 186
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,	
183	Flock, 153
get_parent	Geese, 166
Node, 205	Model< Array_Type, Data_Counter_Type,
get_probabilities	Data_Rule_Type, Data_Rule_Dyn_Type >
Geese, 166	186
get_pset	get_X
Model< Array_Type, Data_Counter_Type,	DEFM, 141
Data_Rule_Type, Data_Rule_Dyn_Type >,	get_X_names
184	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
<pre>get_pset_probs     Model&lt; Array_Type, Data_Counter_Type,</pre>	hasher
Data_Rule_Type, Data_Counter_Type,  Data_Rule_Type, Data_Rule_Dyn_Type >,	counters-meat.hpp, 281, 282
184	hasher_fun
get_pset_stats	Counter< Array_Type, Data_Type >, 133
Model< Array_Type, Data_Counter_Type,	counters-meat.hpp, 282
Data_Rule_Type, Data_Rule_Dyn_Type >,	hasher_fun_
184, 185	counters-meat.hpp, 285
get_rengine	Hasher_fun_type
Geese, 166	typedefs.hpp, 339
Model Array_Type, Data_Counter_Type,	hashes Support< Array_Type, Data_Counter_Type.
Data_Rule_Type, Data_Rule_Dyn_Type >,	Data_Rule_Type, Data_Rule_Dyn_Type >:
185	243
get_row_vec BArray< Cell_Type, Data_Type >, 65	hashes initialized
BArrayDense< Cell Type, Data Type >, 86, 87	Support< Array_Type, Data_Counter_Type,
get_rules	Data_Rule_Type, Data_Rule_Dyn_Type >
Model< Array_Type, Data_Counter_Type,	243
Data_Rule_Type, Data_Rule_Dyn_Type >,	
185	i a a a a a a a a a a a a a a a a a a a
Support< Array_Type, Data_Counter_Type,	counters-meat.hpp, 285
Data_Rule_Type, Data_Rule_Dyn_Type >,	id Node 206
239	Node, 206 idx
get_rules_dyn	Phylo rules, 40, 41
Model < Array_Type, Data_Counter_Type,	if
Data_Rule_Type, Data_Rule_Dyn_Type >, 185	counters-meat.hpp, 282
Support< Array_Type, Data_Counter_Type,	IF_MATCHES
Data_Rule_Type, Data_Rule_Dyn_Type >,	Counting, 15
Data_ridie_type, Data_ridie_Dyfi_type >,	IF_NOTMATCHES

Counting, 15	include/barry/models/geese/geese-node-bones.hpp,
include/barry/barray-bones.hpp, 247	323
include/barry/barray-iterator.hpp, 247	include/barry/models/geese/geese-types.hpp, 323
include/barry/barray-meat-operators.hpp, 248	include/barry/powerset-bones.hpp, 326
include/barry/barray-meat-operators.npp, 240	include/barry/powerset-meat.hpp, 327
	· · · · · · · · · · · · · · · · · · ·
include/barry/barraycell-bones.hpp, 253	include/barry/progress.hpp, 327
include/barry/barraycell-meat.hpp, 254	include/barry/rules-bones.hpp, 328
include/barry/barraydense-bones.hpp, 254	include/barry/rules-meat.hpp, 329
include/barry/barraydense-meat-operators.hpp, 255	include/barry/statscounter-bones.hpp, 329
include/barry/barraydense-meat.hpp, 258	include/barry/statscounter-meat.hpp, 330
include/barry/barraydensecell-bones.hpp, 259	include/barry/support-bones.hpp, 335
include/barry/barraydensecell-meat.hpp, 260	include/barry/support-meat.hpp, 336
include/barry/barraydensecol-bones.hpp, 261	include/barry/typedefs.hpp, 337
include/barry/barraydenserow-bones.hpp, 262	indices
include/barry/barrayrow-bones.hpp, 263	NetCounterData, 200
include/barry/barrayrow-meat.hpp, 263	Phylo rules, 44, 45
include/barry/barrayvector-bones.hpp, 265	inherit_support
include/barry/barrayvector-meat.hpp, 266	Geese, 167
include/barry/barry-configuration.hpp, 266	init
include/barry/barry-debug.hpp, 268	Counter< Array_Type, Data_Type >, 131
include/barry/barry-macros.hpp, 269	DEFM, 142
include/barry/barry.hpp, 270	Flock, 153
include/barry/cell-bones.hpp, 273	Geese, 167
include/barry/cell-meat.hpp, 274	Phylo rules, 45
include/barry/col-bones.hpp, 274	init_fun
include/barry/counters-bones.hpp, 274	Counter< Array_Type, Data_Type >, 133
include/barry/counters-meat.hpp, 275	counters-meat.hpp, 283
include/barry/counters/network-css.hpp, 287	init_fun_
include/barry/counters/network.hpp, 295	counters-meat.hpp, 286
include/barry/freqtable.hpp, 302	init_node
include/barry/model-bones.hpp, 303	Geese, 167
include/barry/model-meat.hpp, 303	init_support
include/barry/models/defm.hpp, 304	PowerSet < Array_Type, Data_Rule_Type >, 218
include/barry/models/defm/counters.hpp, 305	Support< Array_Type, Data_Counter_Type,
include/barry/models/defm/defm-bones.hpp, 310	Data_Rule_Type, Data_Rule_Dyn_Type >,
include/barry/models/defm/defm-meat.hpp, 311	239
include/barry/models/defm/defm-types.hpp, 312	INITIALIZED
include/barry/models/defm/formula.hpp, 313	geese-bones.hpp, 317
include/barry/models/geese.hpp, 315	initialized
include/barry/models/geese/counters.hpp, 308	Flock, 156
include/barry/models/geese/flock-bones.hpp, 315	Geese, 173
include/barry/models/geese/flock-meat.hpp, 316	insert_cell
include/barry/models/geese/geese-bones.hpp, 316	BArray< Cell_Type, Data_Type >, 66
include/barry/models/geese/geese-meat-constructors.hpp	o, BArrayDense < Cell_Type, Data_Type >, 87
318	is_col
include/barry/models/geese/geese-meat-likelihood.hpp,	BArrayVector< Cell_Type, Data_Type >, 115
319	BArrayVector_const< Cell_Type, Data_Type >,
include/barry/models/geese/geese-meat-likelihood exhau	
320	is_dense
include/barry/models/geese/geese-meat-predict.hpp,	BArray< Cell_Type, Data_Type >, 66
320	BArrayDense < Cell_Type, Data_Type >, 87
include/barry/models/geese/geese-meat-predict_exhaust.	
321	Counting, 15
include/barry/models/geese/geese-meat-predict_sim.hpp.	
321	Counting, 15
include/barry/models/geese/geese-meat-simulate.hpp,	is_empty
322	BArray< Cell_Type, Data_Type >, 66
include/barry/models/geese/geese-meat.hpp, 322	BArrayDense< Cell_Type, Data_Type >, 88

is_leaf	MAKE_DUPL_VARS
Node, 205	Counting, 16
is_motif	make_hash
DEFM, 142	FreqTable $<$ T $>$ , 159
Phylo rules, 45	Мар
is_row	barry-configuration.hpp, 268
BArrayVector< Cell_Type, Data_Type >, 115	map_to_state_id
BArrayVector_const< Cell_Type, Data_Type >,	Geese, 173
119	MapVec_type
IS_SPECIATION	typedefs.hpp, 340
Counting, 16	max_num_elements
is_true	Support< Array_Type, Data_Counter_Type,
Phylo rules, 41	Data_Rule_Type, Data_Rule_Dyn_Type >,
iter	244
ConstBArrayRowlter< Cell_Type, Data_Type >,	
128	Model< Array_Type, Data_Counter_Type,
	Data_Rule_Type, Data_Rule_Dyn_Type >,
j	179
counters-meat.hpp, 286	model
statscounter-meat.hpp, 335	Flock, 156
otatossamor moatinpp, soc	
keygen_defm	Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 175
defm-meat.hpp, 312	
keygen_full	~Model, 180
geese-bones.hpp, 317	add_array, 180
keys2support	add_counter, 180, 181
Model< Array_Type, Data_Counter_Type,	add_hasher, 181
Data_Rule_Type, Data_Rule_Dyn_Type >,	add_rule, 181
194	add_rule_dyn, 181, 182
101	arrays2support, 193
lb	colnames, 182
PhyloRuleDynData, 214	conditional_prob, 182
likelihood	counter_fun, 193
Geese, 167	counters, 193
Model< Array_Type, Data_Counter_Type,	delete_counters, 193
Data_Rule_Type, Data_Rule_Dyn_Type >,	delete_rengine, 193
186, 187	delete_rules, 194
likelihood_	delete_rules_dyn, 194
model-meat.hpp, 304	first_calc_done, 194
likelihood_exhaust	gen_key, 183
Geese, 167	get_arrays2support, 183
likelihood_joint	get_counters, 183
Flock, 153	get_norm_const, 183
likelihood total	get_pset, 184
Model< Array_Type, Data_Counter_Type,	get_pset_arrays, 184
Data_Rule_Type, Data_Rule_Dyn_Type >,	get_pset_probs, 184
187	get_pset_stats, 184, 185
logical	get_rengine, 185
-	get_rules, 185
Phylo rules, 45	get_rules_dyn, 185
logodds	get_stats_support, 186
DEFM, 142	get_stats_target, 186
M	get_support_fun, 186
PowerSet < Array_Type, Data_Rule_Type >, 219	keys2support, 194
Support< Array_Type, Data_Counter_Type,	likelihood, 186, 187
Data_Rule_Type, Data_Rule_Dyn_Type >,	likelihood_total, 187
243	Model, 179
MAKE_DEFM_HASHER	normalizing_constants, 194
	nrules, 188
Counting, 16	•

nrules dyn, 188	ncol
nterms, 188	BArray< Cell_Type, Data_Type >, 67
operator=, 188	BArrayDense< Cell_Type, Data_Type >, 88
params_last, 195	Phylo rules, 41
print, 189	NET_C_DATA_IDX
print_stats, 189	network.hpp, 299
pset_arrays, 195	NET C DATA NUM
pset_probs, 195	network.hpp, 299
pset_stats, 195	NetCounter
rengine, 196	network.hpp, 300
-	• •
rules, 196	NetCounterData, 199
rules_dyn, 196	~NetCounterData, 199
sample, 189	indices, 200
set_counters, 190	NetCounterData, 199
set_rengine, 190	numbers, 200
set_rules, 190	NetCounters
set_rules_dyn, 190	network.hpp, 300
set_seed, 191	NetModel
set_transform_model, 191	network.hpp, 301
size, 191	NetRule
size_unique, 192	network.hpp, 301
stats_support, 196	NetRules
stats_support_n_arrays, 197	network.hpp, 301
stats_target, 197	NetStatsCounter
store_psets, 192	network.hpp, 301
support_fun, 197	NetSupport
support_size, 192	network.hpp, 301
transform_model, 192	Network
transform_model_fun, 197	network.hpp, 301
transform_model_term_names, 198	Network counters, 24
with_pset, 198	counter_absdiff, 26
model-meat.hpp	counter_ctriads, 27
likelihood_, 304	counter_degree, 27
update_normalizing_constant, 304	counter_density, 27
motif_census	counter_diff, 28
DEFM, 143	counter_edges, 28
	counter_fixed_effect, 28
N	counter_idegree, 29
PowerSet < Array_Type, Data_Rule_Type >, 220	counter_idegree15, 29
Support< Array_Type, Data_Counter_Type,	counter_isolates, 29, 30
Data_Rule_Type, Data_Rule_Dyn_Type >,	counter_istar2, 30
244	counter_logit_intercept, 30
n_counters	counter_mutual, 30
Support< Array_Type, Data_Counter_Type,	counter_nodecov, 31
Data_Rule_Type, Data_Rule_Dyn_Type >,	counter nodeicov, 31
244	counter_nodematch, 31
n_free	counter_nodeocov, 31
PowerSet< Array_Type, Data_Rule_Type >, 220	counter_odegree, 31, 32
n_locked	counter_odegree15, 32
PowerSet< Array_Type, Data_Rule_Type >, 220	counter_ones, 32
name	counter_ostar2, 33
Counter< Array_Type, Data_Type >, 133	counter_transition, 33
counters-meat.hpp, 283	counter_transition_formula, 34
name_	counter_ttriads, 34
counters-meat.hpp, 286	NETWORK_COUNTER, 34
nannotations	rules_dont_become_zero, 34
Geese, 168	rules_markov_fixed, 35
narray	network-css.hpp
Node, 206	rr

counter_css_census01, 291	NetworkData, 201
counter_css_census02, 291	vertex_attr, 202
counter_css_census03, 291	NetworkDense
counter_css_census04, 292	network.hpp, 302
counter_css_census05, 292	NETWORKDENSE_COUNTER_LAMBDA
counter_css_census06, 292	network.hpp, 300
counter_css_census07, 292	next
counter_css_census08, 293	Progress, 222
counter_css_census09, 293	nfunctions
counter css census10, 293	Flock, 156
counter_css_completely_false_recip_comiss, 293	Geese, 173
counter_css_completely_false_recip_omiss, 294	nfuns
counter_css_mixed_recip, 294	Flock, 153
counter_css_partially_false_recip_commi, 294	Geese, 168
counter_css_partially_false_recip_omiss, 295	nleafs
CSS APPEND, 288	Flock, 154
CSS_CASE_ELSE, 289	Geese, 168
CSS_CASE_PERCEIVED, 289	nnodes
CSS_CASE_TRUTH, 289	Flock, 154
CSS_CHECK_SIZE, 289	Geese, 168
CSS_CHECK_SIZE_INIT, 289	nnozero
CSS_MATCH_TYPE, 290	BArray< Cell_Type, Data_Type >, 67
CSS_NET_COUNTER_LAMBDA_INIT, 290	BArrayDense< Cell_Type, Data_Type >, 88
CSS_PERCEIVED_CELLS, 290	Node, 202
CSS_SIZE, 290	$\sim$ Node, 204
CSS_TRUE_CELLS, 291	annotations, 205
network.hpp	array, 205
BARRY_ZERO_NETWORK, 298	arrays, 206
BARRY_ZERO_NETWORK_DENSE, 298	duplication, 206
NET_C_DATA_IDX, 299	get_parent, 205
NET_C_DATA_NUM, 299	id, 206
NetCounter, 300	is_leaf, 205
NetCounters, 300	narray, 206
NetModel, 301	Node, 203, 204
NetRule, 301	noffspring, 205
NetRules, 301	offspring, 206
NetStatsCounter, 301	ord, 207
NetSupport, 301	parent, 207
Network, 301	probability, 207
NETWORK COUNTER, 299	subtree_prob, 207
NETWORK_COUNTER_LAMBDA, 299	visited, 207
NETWORK_RULE, 299	NodeData, 208
NETWORK_RULE_LAMBDA, 300	blengths, 209
NetworkDense, 302	duplication, 209
NETWORKDENSE_COUNTER_LAMBDA, 300	NodeData, 208
rules_zerodiag, 302	states, 209
NETWORK COUNTER	
<del>_</del>	nodes
Network counters, 34	Geese, 173
network.hpp, 299	noexcept
NETWORK_COUNTER_LAMBDA	counters-meat.hpp, 286
network.hpp, 299	noffspring
NETWORK_RULE	Node, 205
network.hpp, 299	NONE
NETWORK_RULE_LAMBDA	CHECK, 56
network.hpp, 300	EXISTS, 58
NetworkData, 200	normalizing_constants
$\sim$ NetworkData, 202	Model< Array_Type, Data_Counter_Type,
directed, 202	Data_Rule_Type, Data_Rule_Dyn_Type >,

194	BArrayVector_const< Cell_Type, Data_Type >
nrow BArray< Cell_Type, Data_Type >, 67	120
	operator<=  RArrayColl const < Coll Type Data Type > 79
BArrayDense< Cell_Type, Data_Type >, 88 Phylo rules, 41	BArrayCell_const< Cell_Type, Data_Type >, 78 BArrayRow_const< Cell_Type, Data_Type >, 113
nrules	BArrayVector_const< Cell_Type, Data_Type >, 113
Model < Array_Type, Data_Counter_Type,	120
Data_Rule_Type, Data_Rule_Dyn_Type >,	operator>
188	BArrayCell_const< Cell_Type, Data_Type >, 78
nrules_dyn	BArrayRow_const< Cell_Type, Data_Type >, 113
Model< Array_Type, Data_Counter_Type,	BArrayVector_const< Cell_Type, Data_Type >
Data_Rule_Type, Data_Rule_Dyn_Type >,	120
188	operator>=
nterms	BArrayCell_const< Cell_Type, Data_Type >, 78
Flock, 154	BArrayRow_const< Cell_Type, Data_Type >, 113
Geese, 168	BArrayVector_const< Cell_Type, Data_Type >
Model < Array_Type, Data_Counter_Type,	120
Data_Rule_Type, Data_Rule_Dyn_Type >,	operator*=
188	BArray< Cell_Type, Data_Type >, 67
ntrees	BArrayCell< Cell_Type, Data_Type >, 75
Flock, 154	BArrayDense< Cell_Type, Data_Type >, 89
num	BArrayDenseCell< Cell_Type, Data_Type >, 97
Phylo rules, 41, 42	BArrayRow< Cell_Type, Data_Type >, 110
numbers	BArrayVector < Cell_Type, Data_Type >, 116
NetCounterData, 200	operator()
Phylo rules, 45, 46	BArray< Cell_Type, Data_Type >, 67
	barray-meat-operators.hpp, 251
obs_start	BArrayDense< Cell_Type, Data_Type >, 88, 89
Phylo rules, 46	BArrayDenseCol< Cell_Type, Data_Type >, 101
observed_counts	${\tt BArrayDenseCol\_const}{<}{\tt Cell\_Type},{\tt Data\_Type}{>}$
Geese, 169	103
offspring	BArrayDenseRow< Cell_Type, Data_Type >, 106
Node, 206	BArrayDenseRow_const< Cell_Type, Data_Type
ONE	>, 108
CHECK, 56 EXISTS, 58	Flock, 154
operator BArrayRow< Cell_Type, Data_Type >	Phylo rules, 42
BArrayRow< Cell_Type, Data_Type >, 110	PhyloCounterData, 211
operator BArrayRow_const< Cell_Type, Data_Type >	PhyloRuleDynData, 213
BArrayRow_const< Cell_Type, Data_Type >, 112	Rule < Array_Type, Data_Type >, 225
operator Cell_Type	Rules< Array_Type, Data_Type >, 228 vecHasher< T >, 245
BArrayCell< Cell_Type, Data_Type >, 75	operator+=
BArrayCell_const< Cell_Type, Data_Type >, 77	BArray< Cell_Type, Data_Type >, 68
BArrayDenseCell< Cell_Type, Data_Type >, 97	BArrayCell< Cell_Type, Data_Type >, 75
Cell< Cell_Type >, 124	BArrayDense < Cell_Type, Data_Type >, 89
operator std::vector< Cell_Type >	BArrayDenseCell< Cell_Type, Data_Type >, 98
BArrayVector< Cell_Type, Data_Type >, 116	BArrayRow< Cell_Type, Data_Type >, 110
BArrayVector_const< Cell_Type, Data_Type >,	BArrayVector< Cell_Type, Data_Type >, 116
119	operator-=
operator!=	BArray< Cell_Type, Data_Type >, 68
BArrayCell_const< Cell_Type, Data_Type >, 77	BArrayCell< Cell_Type, Data_Type >, 75
BArrayRow_const< Cell_Type, Data_Type >, 112	BArrayDense< Cell_Type, Data_Type >, 90
BArrayVector_const< Cell_Type, Data_Type >,	BArrayDenseCell< Cell_Type, Data_Type >, 98
119	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	BArray< Cell_Type, Data_Type >, 69
BArrayRow_const< Cell_Type, Data_Type >, 112	BArrayCell< Cell_Type, Data_Type >, 75

BArrayDense< Cell_Type, Data_Type >, 90	counter_longest, 50
BArrayDenseCell< Cell_Type, Data_Type >, 98	counter_loss, 51
BArrayRow< Cell_Type, Data_Type >, 111	counter_maxfuns, 51
BArrayVector< Cell_Type, Data_Type >, 116	counter_neofun, 51
operator=	counter_neofun_a2b, 51
BArray< Cell_Type, Data_Type >, 69	counter_overall_changes, 52
BArrayCell< Cell_Type, Data_Type >, 75	counter_overall_gains, 52
BArrayDense < Cell_Type, Data_Type >, 90, 91	counter_overall_gains_from_0, 52
BArrayDenseCell< Cell_Type, Data_Type >, 98	counter_overall_loss, 52
BArrayRow< Cell_Type, Data_Type >, 111	counter_pairwise_first_gain, 53
BArrayVector< Cell_Type, Data_Type >, 117	counter_pairwise_neofun_singlefun, 53
Cell< Cell_Type >, 124, 125	counter_pairwise_overall_change, 53
Counter < Array_Type, Data_Type >, 131, 132	counter_pairwise_preserving, 53
Counters< Array_Type, Data_Type >, 137	counter_preserve_pseudogene, 54
Geese, 169	counter_prop_genes_changing, 54
Model < Array_Type, Data_Counter_Type,	counter_subfun, 54
Data_Rule_Type, Data_Rule_Dyn_Type >,	Phylo rules, 35
188	~DEFMCounterData, 43
Rules < Array_Type, Data_Type >, 229	~DEFMData, 43
operator== BArray< Cell_Type, Data_Type >, 69	~DEFMRuleDynData, 43
BArrayCell< Cell_Type, Data_Type >, 09  BArrayCell< Cell_Type, Data_Type >, 76	array, 44 at, 39
BArrayCell_const< Cell_Type, Data_Type >, 78	counts, 44
BArrayDense < Cell_Type, Data_Type >, 91	covar_sort, 44
BArrayDenseCell< Cell_Type, Data_Type >, 99	covar_used, 44
BArrayRow< Cell_Type, Data_Type >, 111	covariates, 44
BArrayRow_const< Cell_Type, Data_Type >, 113	DEFMCounter, 37
BArrayVector< Cell_Type, Data_Type >, 117	DEFMCounterData, 39
BArrayVector_const< Cell_Type, Data_Type >,	DEFMCounters, 37
120	DEFMData, 39
Cell< Cell_Type >, 125	DEFMModel, 37
operator[]	DEFMRule, 38
Counters< Array_Type, Data_Type >, 138	DEFMRuleData, 40
PhyloCounterData, 211	DEFMRuleDyn, 38
PowerSet < Array_Type, Data_Rule_Type >, 218	DEFMRuleDynData, 40
ord	DEFMRules, 38
Node, 207	DEFMRulesDyn, 38
out_of_range	DEFMStatsCounter, 38
BArray< Cell_Type, Data_Type >, 69	DEFMSupport, 38
BArrayDense < Cell_Type, Data_Type >, 91	idx, 40, 41
narama laat	indices, 44, 45
params_last  Model< Array_Type, Data_Counter_Type,	init, 45
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,	is_motif, 45
195	is_true, 41
parent	logical, 45
Node, 207	ncol, 41
parse_polytomies	nrow, 41
Flock, 155	num, 41, 42
Geese, 169	numbers, 45, 46
Phylo counters, 47	obs_start, 46 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	Counting, 16
counter_k_genes_changing, 50	PHYLO_COUNTER_LAMBDA
counter_less_than_p_prop_genes_changing, 50	

Counting, 17	PhyloRuleDynData, 214
PHYLO_RULE_DYN_LAMBDA	POS N
Counting, 17	barraydense-meat-operators.hpp, 256
PhyloArray	barraydense-meat.hpp, 259
geese-types.hpp, 324	barraydensecol-bones.hpp, 261
	• • • • • • • • • • • • • • • • • • • •
PhyloCounter	barraydenserow-bones.hpp, 263
geese-types.hpp, 324	PowerSet < Array Time Deta Dula Time > 216
PhyloCounterData, 209	PowerSet < Array_Type, Data_Rule_Type >, 216
at, 210	PowerSet < Array_Type, Data_Rule_Type >, 214
begin, 210	~PowerSet, 216
empty, 210	add_rule, 216, 217
end, 211	begin, 217
get_counters, 211	calc, 217
operator(), 211	coordinates_free, 219
operator[], 211	coordinates_locked, 219
PhyloCounterData, 210	data, 219
push_back, 211	EmptyArray, 219
reserve, 211	end, 217
shrink_to_fit, 212	get_data, 217
size, 212	get_data_ptr, 218
PhyloCounters	init_support, 218
geese-types.hpp, 325	M, 219
PhyloModel	N, 220
geese-types.hpp, 325	n_free, 220
PhyloPowerSet	n_locked, 220
geese-types.hpp, 325	operator[], 218
PhyloRule	PowerSet, 216
geese-types.hpp, 325	reset, 218
PhyloRuleData	rules, 220
geese-types.hpp, 325	rules_deleted, 220
PhyloRuleDyn	size, 218
geese-types.hpp, 325	predict
PhyloRuleDynData, 212	Geese, 169
~PhyloRuleDynData, 213	predict_backend
counts, 213	Geese, 170
duplication, 213	predict_exhaust
lb, 214	Geese, 170
operator(), 213	predict_exhaust_backend
PhyloRuleDynData, 213	Geese, 170
pos, 214	predict_sim
ub, 214	Geese, 170
PhyloRules	print
geese-types.hpp, 326	BArray< Cell_Type, Data_Type >, 69
PhyloRulesDyn	BArrayDense< Cell_Type, Data_Type >, 91
geese-types.hpp, 326	DEFM, 143
PhyloStatsCounter	Flock, 155
geese-types.hpp, 326	FreqTable $<$ T $>$ , 159
PhyloSupport	Geese, 170
geese-types.hpp, 326	Model < Array_Type, Data_Counter_Type
POS	Data_Rule_Type, Data_Rule_Dyn_Type >
barraydense-meat-operators.hpp, 256	189
barraydense-meat.hpp, 258	Phylo rules, 42
barraydensecell-bones.hpp, 260	Support< Array_Type, Data_Counter_Type
barraydensecell-meat.hpp, 260	Data_Rule_Type, Data_Rule_Dyn_Type >:
barraydensecel-meathpp, 200 barraydensecol-bones.hpp, 261	240
barraydenserow-bones.hpp, 263	print_n  RArray < Coll Type Data Type > 70
geese-types.hpp, 324	BArray < Cell_Type, Data_Type >, 70
pos	print_nodes

Geese, 171	BArrayDense < Cell_Type, Data_Type >, 92
print_observed_counts	Entries< Cell_Type >, 149
Geese, 171	statscounter-meat.hpp, 332
print_stats	return
Model Array_Type, Data_Counter_Type,	counters-meat.hpp, 287
Data_Rule_Type, Data_Rule_Dyn_Type >,	statscounter-meat.hpp, 335
189	rhs
printf_barry	barray-meat-operators.hpp, 251
barry-configuration.hpp, 267	rm_cell
probability	BArray< Cell_Type, Data_Type >, 70
Node, 207	BArrayDense < Cell_Type, Data_Type >, 92
Progress, 221	ROW
~Progress, 221	barray-meat-operators.hpp, 249
end, 222	barray-meat.hpp, 253
next, 222	barraydense-meat-operators.hpp, 256
Progress, 221	barraydense-meat.hpp, 259
progress.hpp	row
BARRY_PROGRESS_BAR_WIDTH, 328	BArray< Cell_Type, Data_Type >, 70
pset_arrays	BArrayDense < Cell_Type, Data_Type >, 92
Model Array_Type, Data_Counter_Type,	Row_type
Data_Rule_Type, Data_Rule_Dyn_Type >,	typedefs.hpp, 340
195	rowsum
pset_loc	BArrayDense < Cell_Type, Data_Type >, 92 Rule
Geese, 173	
pset_probs	Rule < Array_Type, Data_Type >, 223
Model Array_Type, Data_Counter_Type,	Rule < Array_Type, Data_Type >, 222 ~Rule, 223
Data_Rule_Type, Data_Rule_Dyn_Type >, 195	D, 224
pset_stats  Model< Array_Type, Data_Counter_Type,	get_description, 224
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,	get_name, 224 operator(), 225
195	Rule, 223
push_back	rule_dyn_limit_changes
PhyloCounterData, 211	Phylo rules, 42
Thylodouniol Bala, 211	rule_fun_default
README.md, 342	rules-bones.hpp, 329
reduced_sequence	Rule_fun_type
Geese, 174	typedefs.hpp, 340
rengine	RULE FUNCTION
Flock, 156	barry.hpp, 273
Model< Array_Type, Data_Counter_Type,	geese-bones.hpp, 317
Data_Rule_Type, Data_Rule_Dyn_Type >,	RULE LAMBDA
196	barry.hpp, 273
res	Rules
counters-meat.hpp, 286	Rules< Array_Type, Data_Type >, 226
reserve	rules
BArray< Cell_Type, Data_Type >, 70	Model < Array_Type, Data_Counter_Type
BArrayDense < Cell_Type, Data_Type >, 91	Data_Rule_Type, Data_Rule_Dyn_Type >
FreqTable < T >, 159	196
PhyloCounterData, 211	PowerSet < Array_Type, Data_Rule_Type >, 220
reset	Rules < Array_Type, Data_Type >, 225
PowerSet < Array_Type, Data_Rule_Type >, 218	$\sim$ Rules, 226
reset_array	add_rule, 227
StatsCounter < Array_Type, Data_Type >, 232	begin, 227
Support	end, 227
Data_Rule_Type, Data_Rule_Dyn_Type >,	get_descriptions, 227
240	get_names, 228
resize  PArroy < Coll Type Data Type > 70	get_seq, 228
BArray< Cell_Type, Data_Type >, 70	

operator(), 228	Flock, 155
operator=, 229	Geese, 171
Rules, 226	Model < Array_Type, Data_Counter_Type,
size, 229	Data_Rule_Type, Data_Rule_Dyn_Type >,
rules-bones.hpp	191
rule_fun_default, 329	set_transform_model
rules_deleted	Model< Array_Type, Data_Counter_Type,
PowerSet < Array_Type, Data_Rule_Type >, 220	Data_Rule_Type, Data_Rule_Dyn_Type >,
rules_dont_become_zero	191
Network counters, 34	shrink_to_fit
rules_dyn  Model< Array Type, Data Counter Type,	PhyloCounterData, 212
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,	simulate DEFM, 143
196	Geese, 171
rules_markov_fixed	size
Network counters, 35	BArrayDenseCol< Cell_Type, Data_Type >, 101
rules zerodiag	BArrayDenseCol_const< Cell_Type, Data_Type >,
network.hpp, 302	104
	BArrayDenseRow< Cell_Type, Data_Type >, 106
sample	BArrayDenseRow_const< Cell_Type, Data_Type
Model< Array_Type, Data_Counter_Type,	>, 108
Data_Rule_Type, Data_Rule_Dyn_Type >,	BArrayVector< Cell_Type, Data_Type >, 117
189	BArrayVector_const< Cell_Type, Data_Type >,
sequence	121
Geese, 174	Counters< Array_Type, Data_Type >, 138
set_counters	FreqTable $<$ T $>$ , 160
Model < Array_Type, Data_Counter_Type,	Model< Array_Type, Data_Counter_Type,
Data_Rule_Type, Data_Rule_Dyn_Type >,	$Data_Rule_Type, Data_Rule_Dyn_Type >,$
190 State Country & Arroy Time Date Time > 222	191
StatsCounter < Array_Type, Data_Type >, 233	PhyloCounterData, 212
Support < Array_Type, Data_Counter_Type,	PowerSet < Array_Type, Data_Rule_Type >, 218
Data_Rule_Type, Data_Rule_Dyn_Type >, 240	Rules< Array_Type, Data_Type >, 229
set_data	StatsCounter< Array_Type, Data_Type >, 233
BArray< Cell_Type, Data_Type >, 71	size_unique
BArrayDense < Cell_Type, Data_Type >, 93	Model < Array_Type, Data_Counter_Type,
set_hasher	Data_Rule_Type, Data_Rule_Dyn_Type >,
Counter< Array_Type, Data_Type >, 132	192
set_names	sort_array
DEFM, 143	typedefs.hpp, 340 source
set_rengine	Entries < Cell_Type >, 149
Model< Array_Type, Data_Counter_Type,	states
Data_Rule_Type, Data_Rule_Dyn_Type >,	NodeData, 209
190	Statistical Models, 24
set_rules	stats_support
Model< Array_Type, Data_Counter_Type,	Model< Array_Type, Data_Counter_Type,
Data_Rule_Type, Data_Rule_Dyn_Type >,	Data Rule Type, Data Rule Dyn Type >,
190	196
Support< Array_Type, Data_Counter_Type,	stats_support_n_arrays
Data_Rule_Type, Data_Rule_Dyn_Type >,	Model< Array_Type, Data_Counter_Type,
241	Data_Rule_Type, Data_Rule_Dyn_Type >,
set_rules_dyn	197
Model Array_Type, Data_Counter_Type,	stats_target
Data_Rule_Type, Data_Rule_Dyn_Type >,	Model < Array_Type, Data_Counter_Type,
190 Support Arroy Type Date Counter Type	$Data_Rule_Type, Data_Rule_Dyn_Type >,$
Support < Array_Type, Data_Counter_Type,	197
Data_Rule_Type, Data_Rule_Dyn_Type >, 241	StatsCounter
set seed	StatsCounter< Array_Type, Data_Type >, 230,

231	current_stats, 242
StatsCounter< Array_Type, Data_Type >, 229	delete_counters, 242
∼StatsCounter, 231	delete_rules, 243
add_counter, 231	delete_rules_dyn, 243
count_all, 231	eval_rules_dyn, 238
count current, 232	get_counters, 238
count_init, 232	get_counts, 238
get_counters, 232	get_current_stats, 238
get_descriptions, 232	get data, 239
get_names, 232	get_rules, 239
reset_array, 232	get_rules_dyn, 239
set_counters, 233	hashes, 243
size, 233	hashes_initialized, 243
StatsCounter, 230, 231	init_support, 239
statscounter-meat.hpp	M, 243
clear, 331	max_num_elements, 244
	N, 244
counter, 333	
counter_deleted, 334	n_counters, 244
counters, 334	print, 240
counters_, 334	reset_array, 240
current_stats, 334	set_counters, 240
EmptyArray, 334	set_rules, 241
f_, 335	set_rules_dyn, 241
for, 331	Support, 235, 236
j, 335	support-meat.hpp
resize, 332	BARRY_SUPPORT_MEAT_HPP, 336
return, 335	support_fun
STATSCOUNTER_TEMPLATE, 331–333	Model< Array_Type, Data_Counter_Type,
STATSCOUNTER_TEMPLATE_ARGS, 331	Data_Rule_Type, Data_Rule_Dyn_Type >,
STATSCOUNTER_TYPE, 331	197
STATSCOUNTER_TEMPLATE	support_size
statscounter-meat.hpp, 331–333	Flock, 155
STATSCOUNTER_TEMPLATE_ARGS	Geese, 171
statscounter-meat.hpp, 331	Model < Array_Type, Data_Counter_Type,
STATSCOUNTER_TYPE	Data_Rule_Type, Data_Rule_Dyn_Type >,
statscounter-meat.hpp, 331	192
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
192	swap_cols
subtree_prob	BArray Cell_Type, Data_Type >, 71
Node, 207	BArrayDense< Cell_Type, Data_Type >, 93
Support	swap_rows
Support< Array_Type, Data_Counter_Type,	BArray< Cell_Type, Data_Type >, 71
Data_Rule_Type, Data_Rule_Dyn_Type >,	BArrayDense< Cell_Type, Data_Type >, 93
235, 236	· target
Support < Array_Type, Data_Counter_Type, Data_Rule_T	Entries< Cell_Type >, 149
Data_Rule_Dyn_Type >, 233	this
~Support, 236	barray-meat-operators.hpp, 252
add_counter, 236	TMP_HASHER_CALL
add_rule, 236, 237	counters-meat.hpp, 277
add_rule_dyn, 237	
calc, 237	toggle_cell BArray< Cell_Type, Data_Type >, 72
change_stats, 241	BArrayDense < Cell_Type, Data_Type >, 72  BArrayDense < Cell_Type, Data_Type >, 94
coordiantes_n_free, 241	
coordiantes_n_locked, 242	toggle_lock BArray< Cell_Type, Data_Type >, 72
coordinates_free, 242	
coordinates_locked, 242	BArrayDense < Cell_Type, Data_Type >, 94
	transform_model

Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 192  transform_model_fun Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 197  transform_model_term_names Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198	visited  BArray< Cell_Type, Data_Type >, 73  BArrayDense< Cell_Type, Data_Type >, 96  Cell< Cell_Type >, 125  Node, 207  with_pset  Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 198
transpose	X_ncol
BArray< Cell_Type, Data_Type >, 72	Phylo rules, 46
BArrayDense< Cell_Type, Data_Type >, 94	X_nrow
TWO	Phylo rules, 46
CHECK, 56	7500 0511
EXISTS, 58	ZERO_CELL
typedefs.hpp	barraydense-meat.hpp, 259
Col_type, 339	barraydensecol-bones.hpp, 262
Counter_fun_type, 339	barraydenserow-bones.hpp, 263
Counts type, 339	zero_col
Hasher_fun_type, 339	BArray< Cell_Type, Data_Type >, 72
MapVec_type, 340	BArrayDense < Cell_Type, Data_Type >, 94
Row_type, 340	Zero_row
Rule_fun_type, 340	BArray Cell_Type, Data_Type >, 72
sort_array, 340	BArrayDense < Cell_Type, Data_Type >, 94
vec_equal, 341	
vec_equal_approx, 341	
vec_inner_prod, 341, 342	
ub PhyloRuleDynData, 214 UKNOWN EXISTS, 58 UNI_SUB counters.hpp, 308 update_annotations Geese, 171 update_normalizing_constant model-meat.hpp, 304	
val	
Entries< Cell_Type >, 149	
value	
Cell< Cell_Type >, 125	
vec_diff	
geese-bones.hpp, 317	
vec equal	
typedefs.hpp, 341	
vec_equal_approx	
typedefs.hpp, 341	
vec_inner_prod	
typedefs.hpp, 341, 342	
vecHasher< T >, 244	
operator(), 245	
vector_caster	
geese-bones.hpp, 318	
vertex_attr	
NetworkData, 202	