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 Class Index	7
3.1 Class List	7
4 File Index	9
4.1 File List	9
5 Module Documentation	11
5.1 Counting	11
5.1.1 Detailed Description	11
5.2 Statistical Models	11
5.2.1 Detailed Description	12
5.3 DEFMArray counters	12
5.3.1 Detailed Description	13
5.3.2 Function Documentation	14
5.3.2.1 counter_absdiff()	14
5.3.2.2 counter_ctriads() [1/2]	14
5.3.2.3 counter_ctriads() [2/2]	14
5.3.2.4 counter_degree()	15
5.3.2.5 counter_density()	15
5.3.2.6 counter_diff()	15
5.3.2.7 counter_edges()	15
5.3.2.8 counter_fixed_effect()	15
5.3.2.9 counter_idegree() [1/2]	16
5.3.2.10 counter_idegree() [2/2]	16
5.3.2.11 counter_idegree15() [1/2]	16
5.3.2.12 counter_idegree15() [2/2]	17
5.3.2.13 counter_isolates() [1/2]	17
5.3.2.14 counter_isolates() [2/2]	17
5.3.2.15 counter istar2() [1/2]	17
5.3.2.16 counter_istar2() [2/2]	17
5.3.2.17 counter_mutual()	18
5.3.2.18 counter_nodecov()	18
5.3.2.19 counter_nodeicov()	18
5.3.2.20 counter_nodematch()	18
5.3.2.21 counter_nodeocov()	18
5.3.2.22 counter_odegree() [1/2]	19
5.3.2.23 counter_odegree() [2/2]	19
5.3.2.24 counter_odegree15() [1/2]	19
5.3.2.25 counter_odegree15() [2/2]	19
5.5.2.25 Counter_odegree (5)() [2/2]	13

5.3.2.26 counter_ones()	19
5.3.2.27 counter_ostar2() [1/2]	20
5.3.2.28 counter_ostar2() [2/2]	20
5.3.2.29 counter_transition()	20
5.3.2.30 counter_ttriads() [1/2]	21
5.3.2.31 counter_ttriads() [2/2]	21
5.3.2.32 NETWORK_COUNTER()	21
5.3.2.33 rules_zerodiag()	21
5.4 Phylo counters	22
5.4.1 Detailed Description	23
5.4.2 Function Documentation	23
5.4.2.1 counter_co_opt()	23
5.4.2.2 counter_cogain()	24
5.4.2.3 counter_gains()	24
5.4.2.4 counter_gains_from_0()	24
5.4.2.5 counter_gains_k_offspring()	24
5.4.2.6 counter_genes_changing()	25
5.4.2.7 counter_k_genes_changing()	25
5.4.2.8 counter_less_than_p_prop_genes_changing()	25
5.4.2.9 counter_longest()	25
5.4.2.10 counter_loss()	26
5.4.2.11 counter_maxfuns()	26
5.4.2.12 counter_neofun()	26
5.4.2.13 counter_neofun_a2b()	26
5.4.2.14 counter_overall_changes()	27
5.4.2.15 counter_overall_gains()	27
5.4.2.16 counter_overall_gains_from_0()	27
5.4.2.17 counter_overall_loss()	27
5.4.2.18 counter_pairwise_first_gain()	28
5.4.2.19 counter_pairwise_neofun_singlefun()	28
5.4.2.20 counter_pairwise_overall_change()	28
5.4.2.21 counter_pairwise_preserving()	28
5.4.2.22 counter_preserve_pseudogene()	29
5.4.2.23 counter_prop_genes_changing()	29
5.4.2.24 counter_subfun()	29
5.5 Phylo rules	29
5.5.1 Detailed Description	29
5.5.2 Function Documentation	30
5.5.2.1 rule_dyn_limit_changes()	30
6 Namespace Decumentation	04
6 Namespace Documentation	31
6.1 barry Namespace Reference	31

6.1.1 Detailed Description	. 31
6.2 barry::counters Namespace Reference	. 31
6.2.1 Detailed Description	. 31
6.3 barry::counters::defm Namespace Reference	. 32
6.4 barry::counters::network Namespace Reference	. 32
6.5 barry::counters::phylo Namespace Reference	. 32
6.6 CHECK Namespace Reference	. 32
6.6.1 Detailed Description	. 32
6.6.2 Variable Documentation	. 32
6.6.2.1 BOTH	. 32
6.6.2.2 NONE	. 32
6.6.2.3 ONE	. 33
6.6.2.4 TWO	. 33
6.7 EXISTS Namespace Reference	. 33
6.7.1 Detailed Description	. 33
6.7.2 Variable Documentation	. 33
6.7.2.1 AS_ONE	. 33
6.7.2.2 AS_ZERO	. 34
6.7.2.3 BOTH	. 34
6.7.2.4 NONE	. 34
6.7.2.5 ONE	. 34
6.7.2.6 TWO	. 34
6.7.2.7 UKNOWN	. 34
7 Class Documentation	35
7.1 BArray < Cell_Type, Data_Type > Class Template Reference	
7.1.1 Detailed Description	
7.1.2 Constructor & Destructor Documentation	
7.1.2.1 BArray() [1/6]	
7.1.2.2 BArray() [2/6]	
7.1.2.3 BArray() [3/6]	
7.1.2.4 BArray() [4/6]	
7.1.2.5 BArray() [5/6]	
7.1.2.6 BArray() [6/6]	
7.1.2.7 ~BArray()	
7.1.3 Member Function Documentation	
7.1.3.1 clear()	
7.1.3.2 col()	
7.1.3.3 D() [1/2]	
7.1.3.4 D() [2/2]	
7.1.3.5 D_ptr() [1/2]	
5_pa(/ [±/2]	
7.1.3.6 D_ptr() [2/2]	

7.1.3.7 default_val()
7.1.3.8 flush_data()
7.1.3.9 get_cell()
7.1.3.10 get_col_vec() [1/2]
7.1.3.11 get_col_vec() [2/2]
7.1.3.12 get_entries()
7.1.3.13 get_row_vec() [1/2]
7.1.3.14 get_row_vec() [2/2]
7.1.3.15 insert_cell() [1/3]
7.1.3.16 insert_cell() [2/3]
7.1.3.17 insert_cell() [3/3]
7.1.3.18 is_dense()
7.1.3.19 is_empty()
7.1.3.20 ncol()
7.1.3.21 nnozero()
7.1.3.22 nrow()
7.1.3.23 operator()() [1/2]
7.1.3.24 operator()() [2/2]
7.1.3.25 operator*=()
7.1.3.26 operator+=() [1/3]
7.1.3.27 operator+=() [2/3]
7.1.3.28 operator+=() [3/3]
7.1.3.29 operator-=() [1/3]
7.1.3.30 operator-=() [2/3]
7.1.3.31 operator-=() [3/3]
7.1.3.32 operator/=()
7.1.3.33 operator=() [1/2]
7.1.3.34 operator=() [2/2]
7.1.3.35 operator==()
7.1.3.36 out_of_range()
7.1.3.37 print()
7.1.3.38 reserve()
7.1.3.39 resize()
7.1.3.40 rm_cell()
7.1.3.41 row()
7.1.3.42 set_data()
7.1.3.43 swap_cells()
7.1.3.44 swap_cols()
7.1.3.45 swap_rows()
7.1.3.46 toggle_cell()
7.1.3.47 toggle_lock()
7.1.3.48 transpose()

7.1.3.49 zero_col()	48
7.1.3.50 zero_row()	48
7.1.4 Friends And Related Function Documentation	48
7.1.4.1 BArrayCell< Cell_Type, Data_Type >	48
7.1.4.2 BArrayCell_const< Cell_Type, Data_Type >	49
7.1.5 Member Data Documentation	49
7.1.5.1 visited	49
7.2 BArrayCell< Cell_Type, Data_Type > Class Template Reference	49
7.2.1 Detailed Description	49
7.2.2 Constructor & Destructor Documentation	50
7.2.2.1 BArrayCell()	50
7.2.2.2 ~BArrayCell()	50
7.2.3 Member Function Documentation	50
7.2.3.1 operator Cell_Type()	50
7.2.3.2 operator*=()	50
7.2.3.3 operator+=()	51
7.2.3.4 operator-=()	51
7.2.3.5 operator/=()	51
7.2.3.6 operator=()	51
7.2.3.7 operator==()	51
7.3 BArrayCell_const< Cell_Type, Data_Type > Class Template Reference	52
7.3.1 Detailed Description	52
7.3.2 Constructor & Destructor Documentation	52
7.3.2.1 BArrayCell_const()	52
7.3.2.2 ~BArrayCell_const()	52
7.3.3 Member Function Documentation	53
7.3.3.1 operator Cell_Type()	53
7.3.3.2 operator"!=()	53
7.3.3.3 operator<()	53
7.3.3.4 operator<=()	53
7.3.3.5 operator==()	53
7.3.3.6 operator>()	54
7.3.3.7 operator>=()	54
7.4 BArrayDense< Cell_Type, Data_Type > Class Template Reference	54
7.4.1 Detailed Description	56
7.4.2 Constructor & Destructor Documentation	57
7.4.2.1 BArrayDense() [1/6]	57
7.4.2.2 BArrayDense() [2/6]	57
7.4.2.3 BArrayDense() [3/6]	57
7.4.2.4 BArrayDense() [4/6]	58
7.4.2.5 BArrayDense() [5/6]	58
7.4.2.6 BArrayDense() [6/6]	58

7.4.2.7 ∼l	BArrayDense()	 	 	 	 	58
7.4.3 Member Fur	ction Documentation	 	 	 	 	58
7.4.3.1 cle	ar()	 	 	 	 	58
7.4.3.2 co	() [1/2]	 	 	 	 	59
7.4.3.3 co	() [2/2]	 	 	 	 	59
7.4.3.4 co	sum()	 	 	 	 	59
7.4.3.5 D([1/2]	 	 	 	 	59
7.4.3.6 D([2/2]	 	 	 	 	59
7.4.3.7 D_	ptr() [1/2]	 	 	 	 	59
7.4.3.8 D_	ptr() [2/2]	 	 	 	 	60
7.4.3.9 de	fault_val()	 	 	 	 	60
7.4.3.10 g	et_cell()	 	 	 	 	60
7.4.3.11 g	et_col_vec() [1/2] .	 	 	 	 	60
7.4.3.12 g	et_col_vec() [2/2] .	 	 	 	 	60
7.4.3.13 g	et_data()	 	 	 	 	60
7.4.3.14 g	et_entries()	 	 	 	 	61
7.4.3.15 g	et_row_vec() [1/2]	 	 	 	 	61
7.4.3.16 g	et_row_vec() [2/2]	 	 	 	 	61
7.4.3.17 ir	sert_cell() [1/2]	 	 	 	 	61
7.4.3.18 ir	sert_cell() [2/2]	 	 	 	 	62
7.4.3.19 is	_dense()	 	 	 	 	62
7.4.3.20 is	_empty()	 	 	 	 	62
7.4.3.21 n	col()	 	 	 	 	62
7.4.3.22 n	nozero()	 	 	 	 	62
7.4.3.23 n	row()	 	 	 	 	62
7.4.3.24 o	perator()() [1/2]	 	 	 	 	63
7.4.3.25 o	perator()() [2/2]	 	 	 	 	63
7.4.3.26 o	perator*=()	 	 	 	 	63
7.4.3.27 o	perator+=() [1/3] .	 	 	 	 	63
7.4.3.28 o	perator+=() [2/3] .	 	 	 	 	63
7.4.3.29 o	perator+=() [3/3] .	 	 	 	 	63
7.4.3.30 o	perator-=() [1/3]	 	 	 	 	64
7.4.3.31 o	perator-=() [2/3]	 	 	 	 	64
7.4.3.32 o	perator-=() [3/3]	 	 	 	 	64
7.4.3.33 o	perator/=()	 	 	 	 	64
7.4.3.34 o	perator=() [1/2]	 	 	 	 	64
7.4.3.35 o	perator=() [2/2]	 	 	 	 	64
7.4.3.36 o	perator==()	 	 	 	 	65
7.4.3.37 o	ut_of_range()	 	 	 	 	65
7.4.3.38 p	rint()	 	 	 	 	65
7.4.3.39 re	eserve()	 	 	 	 	65
7.4.3.40 re	esize()	 	 	 	 	65

7.4.3.41 rm_cell()	65
7.4.3.42 row() [1/2]	66
7.4.3.43 row() [2/2]	66
7.4.3.44 rowsum()	66
7.4.3.45 set_data()	66
7.4.3.46 swap_cells()	66
7.4.3.47 swap_cols()	67
7.4.3.48 swap_rows()	67
7.4.3.49 toggle_cell()	67
7.4.3.50 toggle_lock()	67
7.4.3.51 transpose()	68
7.4.3.52 zero_col()	68
7.4.3.53 zero_row()	68
7.4.4 Friends And Related Function Documentation	68
7.4.4.1 BArrayDenseCell< Cell_Type, Data_Type >	68
7.4.4.2 BArrayDenseCol< Cell_Type, Data_Type >	68
7.4.4.3 BArrayDenseCol_const< Cell_Type, Data_Type >	69
7.4.4.4 BArrayDenseRow< Cell_Type, Data_Type >	69
7.4.4.5 BArrayDenseRow_const< Cell_Type, Data_Type >	69
7.4.5 Member Data Documentation	69
7.4.5.1 visited	69
7.5 BArrayDenseCell < Cell_Type, Data_Type > Class Template Reference	69
7.5.1 Detailed Description	70
7.5.2 Constructor & Destructor Documentation	70
7.5.2.1 BArrayDenseCell()	70
7.5.2.2 ~BArrayDenseCell()	70
7.5.3 Member Function Documentation	71
7.5.3.1 operator Cell_Type()	71
7.5.3.2 operator*=()	71
7.5.3.3 operator+=()	71
7.5.3.4 operator-=()	71
7.5.3.5 operator/=()	71
7.5.3.6 operator=()	72
7.5.3.7 operator==()	72
7.5.4 Friends And Related Function Documentation	72
7.5.4.1 BArrayDense < Cell_Type, Data_Type >	72
7.5.4.2 BArrayDenseCol< Cell_Type, Data_Type >	72
7.5.4.3 BArrayDenseCol_const< Cell_Type, Data_Type >	72
7.6 BArrayDenseCell_const< Cell_Type, Data_Type > Class Template Reference	73
7.6.1 Detailed Description	73
7.7 BArrayDenseCol< Cell_Type, Data_Type > Class Template Reference	73
7.7.1 Detailed Description	73

7.7.2 Constructor & Destructor Documentation	73
7.7.2.1 BArrayDenseCol()	74
7.7.3 Member Function Documentation	74
7.7.3.1 begin()	74
7.7.3.2 end()	74
7.7.3.3 operator()()	74
7.7.3.4 size()	74
7.7.4 Friends And Related Function Documentation	75
7.7.4.1 BArrayDense < Cell_Type, Data_Type >	75
7.7.4.2 BArrayDenseCell< Cell_Type, Data_Type >	75
7.7.4.3 BArrayDenseCell_const< Cell_Type, Data_Type >	75
7.8 BArrayDenseCol_const< Cell_Type, Data_Type > Class Template Reference	75
7.8.1 Detailed Description	76
7.8.2 Constructor & Destructor Documentation	76
7.8.2.1 BArrayDenseCol_const()	76
7.8.3 Member Function Documentation	76
7.8.3.1 begin()	76
7.8.3.2 end()	76
7.8.3.3 operator()()	77
7.8.3.4 size()	77
7.8.4 Friends And Related Function Documentation	77
7.8.4.1 BArrayDenseCell< Cell_Type, Data_Type >	77
7.8.4.2 BArrayDenseCell_const< Cell_Type, Data_Type >	77
7.9 BArrayDenseRow< Cell_Type, Data_Type > Class Template Reference	77
7.9.1 Detailed Description	78
7.9.2 Constructor & Destructor Documentation	78
7.9.2.1 BArrayDenseRow()	78
7.9.3 Member Function Documentation	78
7.9.3.1 begin()	78
7.9.3.2 end()	79
7.9.3.3 operator()()	79
7.9.3.4 size()	79
7.9.4 Friends And Related Function Documentation	79
7.9.4.1 BArrayDense < Cell_Type, Data_Type >	79
7.9.4.2 BArrayDenseCell< Cell_Type, Data_Type >	79
7.9.4.3 BArrayDenseCell_const< Cell_Type, Data_Type >	80
7.10 BArrayDenseRow_const< Cell_Type, Data_Type > Class Template Reference	80
7.10.1 Detailed Description	80
7.10.2 Constructor & Destructor Documentation	80
7.10.2.1 BArrayDenseRow_const()	81
7.10.3 Member Function Documentation	81
7.10.3.1 begin()	81

81
81
81
82
82
82
82
82
83
83
83
83
83
83
83
84
84
84
84
84
85
85
85
85
85
85
85
86
86
86
86
86
86
87
87
87
88
88
88
88
88
89

7.13.3.5 operator std::vector< Cell_Type >()	89
7.13.3.6 operator*=()	89
7.13.3.7 operator+=()	89
7.13.3.8 operator-=()	89
7.13.3.9 operator/=()	90
7.13.3.10 operator=()	90
7.13.3.11 operator==()	90
7.13.3.12 size()	90
7.14 BArrayVector_const< Cell_Type, Data_Type > Class Template Reference	90
7.14.1 Detailed Description	91
7.14.2 Constructor & Destructor Documentation	91
7.14.2.1 BArrayVector_const()	91
7.14.2.2 ~BArrayVector_const()	91
7.14.3 Member Function Documentation	92
7.14.3.1 begin()	92
7.14.3.2 end()	92
7.14.3.3 is_col()	92
7.14.3.4 is_row()	92
7.14.3.5 operator std::vector< Cell_Type >()	92
7.14.3.6 operator"!=()	93
7.14.3.7 operator<()	93
7.14.3.8 operator<=()	93
7.14.3.9 operator==()	93
7.14.3.10 operator>()	93
7.14.3.11 operator>=()	94
7.14.3.12 size()	94
7.15 Cell< Cell_Type > Class Template Reference	94
7.15.1 Detailed Description	95
7.15.2 Constructor & Destructor Documentation	95
7.15.2.1 Cell() [1/7]	95
7.15.2.2 Cell() [2/7]	95
7.15.2.3 ~Cell()	95
7.15.2.4 Cell() [3/7]	96
7.15.2.5 Cell() [4/7]	96
7.15.2.6 Cell() [5/7]	96
7.15.2.7 Cell() [6/7]	96
7.15.2.8 Cell() [7/7]	96
7.15.3 Member Function Documentation	96
7.15.3.1 add() [1/4]	97
7.15.3.2 add() [2/4]	97
7.15.3.3 add() [3/4]	97
7.15.3.4 add() [4/4]	97

7.15.3.5 operator Cell_Type()	7
7.15.3.6 operator"!=()	7
7.15.3.7 operator=() [1/2] 9	8
7.15.3.8 operator=() [2/2] 9	8
7.15.3.9 operator==()	8
7.15.4 Member Data Documentation	8
7.15.4.1 active	8
7.15.4.2 value	8
7.15.4.3 visited	19
7.16 Cell_const< Cell_Type > Class Template Reference	19
7.16.1 Detailed Description	19
7.17 ConstBArrayRowIter< Cell_Type, Data_Type > Class Template Reference	19
7.17.1 Detailed Description	0
7.17.2 Constructor & Destructor Documentation	0
7.17.2.1 ConstBArrayRowlter()	0
7.17.2.2 ~ConstBArrayRowIter()	0
7.17.3 Member Data Documentation	0
7.17.3.1 Array	1
7.17.3.2 current_col	1
7.17.3.3 current_row	1
7.17.3.4 iter	1
7.18 Counter< Array_Type, Data_Type > Class Template Reference	1
7.18.1 Detailed Description	12
7.18.2 Constructor & Destructor Documentation	12
7.18.2.1 Counter() [1/4]	3
7.18.2.2 Counter() [2/4]	3
7.18.2.3 Counter() [3/4]	3
7.18.2.4 Counter() [4/4]	3
7.18.2.5 ~Counter()	3
7.18.3 Member Function Documentation	14
7.18.3.1 count()	14
7.18.3.2 get_description()	14
7.18.3.3 get_name()	14
7.18.3.4 init()	14
7.18.3.5 operator=() [1/2]	14
7.18.3.6 operator=() [2/2]	15
7.18.4 Member Data Documentation	15
7.18.4.1 count_fun	15
7.18.4.2 data	15
7.18.4.3 desc	5
7.18.4.4 init_fun	15
7.18.4.5 name	16

7.19 Counters< Array_Type, Data_Type > Class Template Reference	106
7.19.1 Detailed Description	106
7.19.2 Constructor & Destructor Documentation	107
7.19.2.1 Counters() [1/3]	107
7.19.2.2 ~Counters()	107
7.19.2.3 Counters() [2/3]	107
7.19.2.4 Counters() [3/3]	107
7.19.3 Member Function Documentation	108
7.19.3.1 add_counter() [1/2]	108
7.19.3.2 add_counter() [2/2]	108
7.19.3.3 get_descriptions()	108
7.19.3.4 get_names()	108
7.19.3.5 operator=() [1/2]	108
7.19.3.6 operator=() [2/2]	109
7.19.3.7 operator[]()	109
7.19.3.8 size()	110
7.20 DEFMCounterData Class Reference	110
7.20.1 Detailed Description	110
7.20.2 Constructor & Destructor Documentation	110
7.20.2.1 DEFMCounterData() [1/2]	111
7.20.2.2 DEFMCounterData() [2/2]	111
7.20.2.3 ~DEFMCounterData()	111
7.20.3 Member Function Documentation	111
7.20.3.1 idx()	111
7.20.3.2 num()	111
7.20.4 Member Data Documentation	111
7.20.4.1 indices	112
7.20.4.2 markov_order	112
7.20.4.3 numbers	112
7.21 DEFMData Class Reference	112
7.21.1 Detailed Description	113
7.21.2 Constructor & Destructor Documentation	113
7.21.2.1 DEFMData() [1/2]	113
7.21.2.2 DEFMData() [2/2]	113
7.21.2.3 ~DEFMData()	114
7.21.3 Member Function Documentation	114
7.21.3.1 at()	114
7.21.3.2 operator()()	114
7.21.4 Member Data Documentation	114
7.21.4.1 covariates	115
7.21.4.2 n_covariates	115
7.21.4.3 obs_n_times	115

7.21.4.4 obs_start	115
7.22 Entries < Cell_Type > Class Template Reference	115
7.22.1 Detailed Description	116
7.22.2 Constructor & Destructor Documentation	116
7.22.2.1 Entries() [1/2]	116
7.22.2.2 Entries() [2/2]	116
7.22.2.3 ~Entries()	117
7.22.3 Member Function Documentation	117
7.22.3.1 resize()	117
7.22.4 Member Data Documentation	117
7.22.4.1 source	117
7.22.4.2 target	117
7.22.4.3 val	117
7.23 Flock Class Reference	118
7.23.1 Detailed Description	119
7.23.2 Constructor & Destructor Documentation	119
7.23.2.1 Flock()	119
7.23.2.2 ~Flock()	119
7.23.3 Member Function Documentation	119
7.23.3.1 add_data()	119
7.23.3.2 colnames()	120
7.23.3.3 get_counters()	120
7.23.3.4 get_model()	120
7.23.3.5 get_stats_support()	120
7.23.3.6 get_stats_target()	120
7.23.3.7 get_support_fun()	121
7.23.3.8 init()	121
7.23.3.9 likelihood_joint()	121
7.23.3.10 nfuns()	121
7.23.3.11 nleafs()	122
7.23.3.12 nnodes()	122
7.23.3.13 nterms()	122
7.23.3.14 ntrees()	122
7.23.3.15 operator()()	122
7.23.3.16 parse_polytomies()	123
7.23.3.17 print()	123
7.23.3.18 set_seed()	123
7.23.3.19 support_size()	123
7.23.4 Member Data Documentation	123
7.23.4.1 dat	124
7.23.4.2 initialized	124
7.23.4.3 model	124

7.23.4.4 nfunctions	124
7.23.4.5 rengine	124
7.24 FreqTable < T > Class Template Reference	124
7.24.1 Detailed Description	125
7.24.2 Constructor & Destructor Documentation	125
7.24.2.1 FreqTable()	125
7.24.2.2 ∼FreqTable()	126
7.24.3 Member Function Documentation	126
7.24.3.1 add()	126
7.24.3.2 as_vector()	126
7.24.3.3 clear()	126
7.24.3.4 get_data()	126
7.24.3.5 get_index()	127
7.24.3.6 make_hash()	127
7.24.3.7 print()	127
7.24.3.8 reserve()	127
7.24.3.9 size()	127
7.25 Geese Class Reference	128
7.25.1 Detailed Description	131
7.25.2 Constructor & Destructor Documentation	131
7.25.2.1 Geese() [1/8]	131
7.25.2.2 Geese() [2/8]	132
7.25.2.3 Geese() [3/8]	132
7.25.2.4 Geese() [4/8]	132
7.25.2.5 ~Geese() [1/2]	132
7.25.2.6 Geese() [5/8]	132
7.25.2.7 Geese() [6/8]	133
7.25.2.8 Geese() [7/8]	133
7.25.2.9 Geese() [8/8]	133
7.25.2.10 ~Geese() [2/2]	133
7.25.3 Member Function Documentation	133
7.25.3.1 calc_reduced_sequence() [1/2]	133
7.25.3.2 calc_reduced_sequence() [2/2]	133
7.25.3.3 calc_sequence() [1/2]	134
7.25.3.4 calc_sequence() [2/2]	134
7.25.3.5 colnames() [1/2]	134
7.25.3.6 colnames() [2/2]	134
7.25.3.7 get_annotated_nodes() [1/2]	134
7.25.3.8 get_annotated_nodes() [2/2]	134
7.25.3.9 get_counters() [1/2]	135
7.25.3.10 get_counters() [2/2]	135
7.25.3.11 get_model() [1/2]	135

7.25.3.12 get_model() [2/2]
7.25.3.13 get_probabilities() [1/2]
7.25.3.14 get_probabilities() [2/2]
7.25.3.15 get_rengine() [1/2]
7.25.3.16 get_rengine() [2/2]
7.25.3.17 get_states() [1/2]
7.25.3.18 get_states() [2/2]
7.25.3.19 get_support_fun() [1/2]
7.25.3.20 get_support_fun() [2/2]
7.25.3.21 inherit_support() [1/2]
7.25.3.22 inherit_support() [2/2]
7.25.3.23 init() [1/2]
7.25.3.24 init() [2/2]
7.25.3.25 init_node() [1/2]
7.25.3.26 init_node() [2/2]
7.25.3.27 likelihood() [1/2]
7.25.3.28 likelihood() [2/2]
7.25.3.29 likelihood_exhaust() [1/2]
7.25.3.30 likelihood_exhaust() [2/2]
7.25.3.31 nannotations() [1/2]
7.25.3.32 nannotations() [2/2]
7.25.3.33 nfuns() [1/2]
7.25.3.34 nfuns() [2/2]
7.25.3.35 nleafs() [1/2]
7.25.3.36 nleafs() [2/2]
7.25.3.37 nnodes() [1/2]
7.25.3.38 nnodes() [2/2]
7.25.3.39 nterms() [1/2]
7.25.3.40 nterms() [2/2]
7.25.3.41 observed_counts() [1/2]
7.25.3.42 observed_counts() [2/2]
7.25.3.43 operator=() [1/4]
7.25.3.44 operator=() [2/4]
7.25.3.45 operator=() [3/4]
7.25.3.46 operator=() [4/4]
7.25.3.47 parse_polytomies() [1/2]
7.25.3.48 parse_polytomies() [2/2]
7.25.3.49 predict() [1/2]
7.25.3.50 predict() [2/2]
7.25.3.51 predict_backend() [1/2]
7.25.3.52 predict_backend() [2/2]
7.25.3.53 predict_exhaust() [1/2]

7.25.3.55 predict_exhaust_backend() [1/2]	7.25.3.54 predict_exhaust() [2/2]		142
7.25.3.57 predict_sim() [1/2] 143 7.25.3.58 predict_sim() [2/2] 143 7.25.3.59 print() [1/2] 143 7.25.3.69 print() [2/2] 144 7.25.3.61 print_observed_counts() [1/2] 144 7.25.3.62 print_observed_counts() [2/2] 144 7.25.3.63 set_seed() [1/2] 144 7.25.3.63 set_seed() [1/2] 144 7.25.3.64 set_seed() [2/2] 144 7.25.3.65 simulate() [1/2] 144 7.25.3.66 simulate() [1/2] 145 7.25.3.67 support_size() [1/2] 145 7.25.3.69 support_size() [1/2] 145 7.25.3.69 support_size() [1/2] 145 7.25.3.70 update_annotations() [1/2] 145 7.25.3.70 update_annotations() [1/2] 145 7.25.4.1 delete_rengine 146 7.25.4.2 delete_support 146 7.25.4.3 initialized 146 7.25.4.5 nfunctions 146 7.25.4.5 nindections 146 7.25.4.5 nindections 146 7.25.4.5 nedes 146 7.25.4.6 nodes 146 7.25.4.7 sequence 147 7.25.4.8 reduced_sequence 147 7.25.4.9 sequence 147 7.26.2.1 Model() [1/3] 150 7.26.2.2 Model() [2/3] 151 7.26.2.3 Model() [1/3] 150 7.26.2.2 Model() [1/3] 150 7.26.2.2 Model() [1/3] 151 7.26.2.3 Model() [1/3] 151 7.26.2.3 Model() [1/3] 151 7.26.2.3 Model() [1/3] 151 7.26.2.3 Model() [1/3] 151 7.26.3.3 add_counter() [1/2] 152 7.26.3.4 add_rule() [1/2] 152	7.25.3.55 predict_exhaust_backend() [1/2]		143
7.25.3.58 predict_sim() [2/2] 143 7.25.3.59 print() [1/2] 143 7.25.3.69 print() [1/2] 144 7.25.3.61 print_observed_counts() [1/2] 144 7.25.3.62 print_observed_counts() [2/2] 144 7.25.3.63 set_seed() [1/2] 144 7.25.3.63 set_seed() [1/2] 144 7.25.3.65 simulate() [1/2] 144 7.25.3.65 simulate() [1/2] 144 7.25.3.65 simulate() [1/2] 145 7.25.3.67 support_size() [1/2] 145 7.25.3.68 support_size() [1/2] 145 7.25.3.69 update_annotations() [1/2] 145 7.25.3.70 update_annotations() [1/2] 145 7.25.3.70 update_annotations() [1/2] 145 7.25.4.1 delete_rengine 146 7.25.4.2 delete_support 146 7.25.4.3 initialized 146 7.25.4.5 nfunctions 146 7.25.4.5 nfunctions 146 7.25.4.5 nfunctions 146 7.25.4.5 pset_loc 147 7.25.4.8 reduced_sequence 147 7.25.4.9 sequence 147 7.25.4.9 sequence 147 7.25.4.9 tode() [1/3] 150 7.26.2.1 Model() [1/3] 150 7.26.2.2 Model() [2/3] 151 7.26.2.2 Model() [2/3] 151 7.26.2.3 Model() [1/3] 150 7.26.2.2 Model() [2/3] 151 7.26.2.3 Model() [3/3] 151 7.26.3.3 add_counter() [1/2] 152 7.26.3.3 add_counter() [1/2] 152 7.26.3.3 add_counter() [1/2] 152 7.26.3.4 add_rule() [1/2] 152 7.26.3.4 add_rule() [1/2] 152 7.26.3.5 add_rule() [1/2] 152	7.25.3.56 predict_exhaust_backend() [2/2]		143
7.25.3.59 print[) [1/2] 143 7.25.3.60 print[) [2/2] 144 7.25.3.61 print_observed_counts[) [1/2] 144 7.25.3.62 print_observed_counts[) [2/2] 144 7.25.3.63 set_seed[) [1/2] 144 7.25.3.63 set_seed[) [1/2] 144 7.25.3.64 set_seed[) [2/2] 144 7.25.3.65 simulate[) [1/2] 144 7.25.3.66 simulate[) [1/2] 144 7.25.3.68 simulate[) [1/2] 145 7.25.3.69 support_size[) [1/2] 145 7.25.3.69 support_size[) [1/2] 145 7.25.3.70 update_annotations[) [1/2] 145 7.25.3.70 update_annotations[) [1/2] 145 7.25.3.70 update_annotations[) [1/2] 145 7.25.4.1 delete_rengine 146 7.25.4.2 delete_support 146 7.25.4.3 initialized 146 7.25.4.5 intunctions 146 7.25.4.5 indunctions 146 7.25.4.5 indunctions 146 7.25.4.5 indunctions 146 7.25.4.5 pset_loc 147 7.25.4.9 sequence 147 7.25.4.9 sequence 147 7.25.4.9 sequence 147 7.25.1 Detailed Description 150 7.26.2 Constructor & Destructor Documentation 150 7.26.2 Constructor & Destructor Documentation 150 7.26.2.1 Model() [1/3] 151 7.26.2.2 Model() [1/3] 151 7.26.2.2 Model() [1/3] 151 7.26.2.3 Model() [1/3] 151 7.26.2.4 Model() [1/3] 151 7.26.2.4 Model() [1/3] 151 7.26.2.3 dod_counter() [1/2] 152 7.26.3.3 add_counter() [1/2] 152 7.26.3.4 add_counter() [1/2] 152 7.26.3.5 add_counter() [1/2] 152 7.26.3.5 add_counter() [1/2] 152	7.25.3.57 predict_sim() [1/2]		143
7.25.3.60 print() [2/2] 144 7.25.3.61 print_observed_counts() [1/2] 144 7.25.3.63 set_seed() [1/2] 144 7.25.3.63 set_seed() [1/2] 144 7.25.3.64 set_seed() [2/2] 144 7.25.3.65 simulate() [1/2] 144 7.25.3.65 simulate() [1/2] 144 7.25.3.65 simulate() [2/2] 145 7.25.3.69 support_size() [2/2] 145 7.25.3.69 support_size() [2/2] 145 7.25.3.69 update_annotations() [1/2] 145 7.25.3.70 update_annotations() [1/2] 145 7.25.3.70 update_annotations() [2/2] 145 7.25.4 Member Data Documentation 145 7.25.4.1 delete_rengine 146 7.25.4.2 delete_support 146 7.25.4.3 initialized 146 7.25.4.5 intunctions 146 7.25.4.5 indunctions 146 7.25.4.5 nodes 146 7.25.4.7 pset_loc 147 7.25.4.8 reduced_sequence 147 7.25.4.9 sequence 147 7.25.4.9 sequence 147 7.25.4.9 sequence 147 7.25.1. Detailed Description 150 7.26.2 Constructor & Destructor Documentation 150 7.26.2.1 Model() [1/3] 151 7.26.2.2 Model() [2/3] 151 7.26.2.2 Model() [1/3] 151 7.26.2.3 Member Function Documentation 151 7.26.3.3 add_counter() [1/2] 152 7.26.3.4 add_counter() [1/2] 152 7.26.3.4 add_counter() [1/2] 152 7.26.3.4 add_counter() [1/2] 152 7.26.3.4 add_counter() [1/2] 152 7.26.3.5 add_rule() [1/2] 152	7.25.3.58 predict_sim() [2/2]		143
7.25.3.61 print_observed_counts() [1/2]	7.25.3.59 print() [1/2]		143
7.25.3.62 print_observed_counts() [2/2]	7.25.3.60 print() [2/2]		144
7.25.3.63 set_seed() [1/2]	7.25.3.61 print_observed_counts() [1/2]		144
7.25.3.64 set_seed() [2/2]	7.25.3.62 print_observed_counts() [2/2]		144
7.25.3.65 simulate() [1/2]	7.25.3.63 set_seed() [1/2]		144
7.25.3.66 simulate() [2/2] 145 7.25.3.67 support_size() [1/2] 145 7.25.3.68 support_size() [2/2] 145 7.25.3.68 support_size() [2/2] 145 7.25.3.69 update_annotations() [1/2] 145 7.25.3.70 update_annotations() [2/2] 145 7.25.4 Member Data Documentation 145 7.25.4.1 delete_rengine 146 7.25.4.2 delete_support 146 7.25.4.3 initialized 146 7.25.4.3 initialized 146 7.25.4.4 map_to_nodes 146 7.25.4.5 nfunctions 146 7.25.4.5 nfunctions 146 7.25.4.8 reduced_sequence 147 7.25.4.8 reduced_sequence 147 7.25.4.8 reduced_sequence 147 7.26.1 Detailed Description 150 7.26.2 Constructor & Destructor Documentation 150 7.26.2.2 Model() [1/3] 151 7.26.2.3 Model() [1/3] 151 7.26.3.3 Member Function Documentation 151 7.26.3 Member Function Documentation 151 7.26.3.1 add_array() 151 7.26.3.3 add_counter() [1/2] 152 7.26.3.4 add_rule() [1/2] 152 7.26.3.5 add_rule() [1/2] 152 7.26.3.5 add_rule() [1/2] 152 7.26.3.5 add_rule() [1/2] 152 7.26.3.5 add_rule() [1/2] 152	7.25.3.64 set_seed() [2/2]		144
7.25.3.67 support_size() [1/2]	7.25.3.65 simulate() [1/2]		144
7.25.3.68 support size() [2/2]	7.25.3.66 simulate() [2/2]		145
7.25.3.69 update_annotations() [1/2]	7.25.3.67 support_size() [1/2]		145
7.25.3.70 update_annotations() [2/2]	7.25.3.68 support_size() [2/2]		145
7.25.4 Member Data Documentation 145 7.25.4.1 delete_rengine 146 7.25.4.2 delete_support 146 7.25.4.3 initialized 146 7.25.4.4 map_to_nodes 146 7.25.4.5 nfunctions 146 7.25.4.6 nodes 146 7.25.4.7 pset_loc 147 7.25.4.8 reduced_sequence 147 7.25.4.9 sequence 147 7.26.1 Detailed Description 150 7.26.1 Detailed Description 150 7.26.2 Constructor & Destructor Documentation 150 7.26.2.3 Model() [1/3] 151 7.26.2.3 Model() [1/3] 151 7.26.2.3 Model() [1/3] 151 7.26.3 Member Function Documentation 151 7.26.3 Member Function Documentation 151 7.26.3.1 add_array() 151 7.26.3.2 add_counter() [1/2] 152 7.26.3.3 add_counter() [1/2] 152 7.26.3.3 add_counter() [1/2] 152 7.26.3.4 add_rule() [1/2] 152 7.26.3.5 add_rule() [1/2] 152 7.26.3.5 add_rule() [1/2] 152 7.26.3.5 add_rule() [1/2] 152	7.25.3.69 update_annotations() [1/2]		145
7.25.4.1 delete_rengine	7.25.3.70 update_annotations() [2/2]		145
7.25.4.2 delete_support	7.25.4 Member Data Documentation		145
7.25.4.3 initialized	7.25.4.1 delete_rengine		146
7.25.4.4 map_to_nodes	7.25.4.2 delete_support		146
7.25.4.5 nfunctions	7.25.4.3 initialized		146
7.25.4.6 nodes	7.25.4.4 map_to_nodes		146
7.25.4.7 pset_loc	7.25.4.5 nfunctions		146
7.25.4.8 reduced_sequence	7.25.4.6 nodes		146
7.25.4.9 sequence	7.25.4.7 pset_loc		147
7.26 Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class Template Reference	7.25.4.8 reduced_sequence		147
plate Reference 147 7.26.1 Detailed Description 150 7.26.2 Constructor & Destructor Documentation 150 7.26.2.1 Model() [1/3] 150 7.26.2.2 Model() [2/3] 151 7.26.2.3 Model() [3/3] 151 7.26.2.4 ~ Model() 151 7.26.3 Member Function Documentation 151 7.26.3.1 add_array() 151 7.26.3.2 add_counter() [1/2] 152 7.26.3.3 add_counter() [2/2] 152 7.26.3.4 add_rule() [1/2] 152 7.26.3.5 add_rule() [2/2] 152	7.25.4.9 sequence		147
7.26.1 Detailed Description 150 7.26.2 Constructor & Destructor Documentation 150 7.26.2.1 Model() [1/3] 150 7.26.2.2 Model() [2/3] 151 7.26.2.3 Model() [3/3] 151 7.26.2.4 ~Model() 151 7.26.3 Member Function Documentation 151 7.26.3.1 add_array() 151 7.26.3.2 add_counter() [1/2] 152 7.26.3.3 add_counter() [2/2] 152 7.26.3.4 add_rule() [1/2] 152 7.26.3.5 add_rule() [1/2] 152	7.26 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class	Tem-	
7.26.2 Constructor & Destructor Documentation 150 7.26.2.1 Model() [1/3] 150 7.26.2.2 Model() [2/3] 151 7.26.2.3 Model() [3/3] 151 7.26.2.4 ~Model() 151 7.26.3 Member Function Documentation 151 7.26.3.1 add_array() 151 7.26.3.2 add_counter() [1/2] 152 7.26.3.3 add_counter() [2/2] 152 7.26.3.4 add_rule() [1/2] 152 7.26.3.5 add_rule() [2/2] 152			
7.26.2.1 Model() [1/3]	•		
7.26.2.2 Model() [2/3]	7.26.2 Constructor & Destructor Documentation		150
7.26.2.3 Model() [3/3]	v		
7.26.2.4 ~Model()	7.26.2.2 Model() [2/3]		151
7.26.3 Member Function Documentation 151 7.26.3.1 add_array() 151 7.26.3.2 add_counter() [1/2] 152 7.26.3.3 add_counter() [2/2] 152 7.26.3.4 add_rule() [1/2] 152 7.26.3.5 add_rule() [2/2] 152	7.26.2.3 Model() [3/3]		151
7.26.3.1 add_array()	7.26.2.4 ~Model()		151
7.26.3.2 add_counter() [1/2]	7.26.3 Member Function Documentation		151
7.26.3.3 add_counter() [2/2]	7.26.3.1 add_array()		151
7.26.3.4 add_rule() [1/2]	7.26.3.2 add_counter() [1/2]		152
7.26.3.5 add_rule() [2/2]	7.26.3.3 add_counter() [2/2]		152
	7.26.3.4 add_rule() [1/2]		152
7.26.3.6 add_rule_dyn() [1/2]	7.26.3.5 add_rule() [2/2]		152
	7.26.3.6 add_rule_dyn() [1/2]		152

7.26.3.7 add_rule_dyn() [2/2]	53
7.26.3.8 colnames()	53
7.26.3.9 conditional_prob()	53
7.26.3.10 gen_key()	54
7.26.3.11 get_arrays2support()	54
7.26.3.12 get_counters()	54
7.26.3.13 get_norm_const()	54
7.26.3.14 get_pset()	54
7.26.3.15 get_pset_arrays()	55
7.26.3.16 get_pset_probs()	55
7.26.3.17 get_pset_stats() [1/2]	55
7.26.3.18 get_pset_stats() [2/2]	55
7.26.3.19 get_rengine()	55
7.26.3.20 get_rules()	56
7.26.3.21 get_rules_dyn()	56
7.26.3.22 get_stats_support()	56
7.26.3.23 get_stats_target()	56
7.26.3.24 get_support_fun()	56
7.26.3.25 likelihood() [1/4]	157
7.26.3.26 likelihood() [2/4]	157
7.26.3.27 likelihood() [3/4]	157
7.26.3.28 likelihood() [4/4]	157
7.26.3.29 likelihood_total()	58
7.26.3.30 nterms()	158
7.26.3.31 operator=()	58
7.26.3.32 print()	158
7.26.3.33 print_stats()	58
7.26.3.34 sample() [1/2]	159
7.26.3.35 sample() [2/2]	159
7.26.3.36 set_counters()	59
7.26.3.37 set_keygen()	159
7.26.3.38 set_rengine()	59
7.26.3.39 set_rules()	60
7.26.3.40 set_rules_dyn()	60
7.26.3.41 set_seed()	60
7.26.3.42 set_transform_model()	60
7.26.3.43 size()	61
7.26.3.44 size_unique()	61
7.26.3.45 store_psets()	61
7.26.3.46 support_size()	61
7.26.3.47 transform_model()	61
7.27 NetCounterData Class Reference	62

7.27.1 Detailed Description	62
7.27.2 Constructor & Destructor Documentation	62
7.27.2.1 NetCounterData() [1/2]	62
7.27.2.2 NetCounterData() [2/2]	62
7.27.2.3 ~NetCounterData()	63
7.27.3 Member Data Documentation	63
7.27.3.1 indices	63
7.27.3.2 numbers	63
7.28 NetworkData Class Reference	63
7.28.1 Detailed Description	64
7.28.2 Constructor & Destructor Documentation	64
7.28.2.1 NetworkData() [1/3]	64
7.28.2.2 NetworkData() [2/3]	64
7.28.2.3 NetworkData() [3/3]	64
7.28.2.4 ~NetworkData()	65
7.28.3 Member Data Documentation	65
7.28.3.1 directed	65
7.28.3.2 vertex_attr	65
7.29 Node Class Reference	65
7.29.1 Detailed Description	66
7.29.2 Constructor & Destructor Documentation	66
7.29.2.1 Node() [1/5] 1	67
7.29.2.2 Node() [2/5] 1	67
7.29.2.3 Node() [3/5]	67
7.29.2.4 Node() [4/5] 1	67
7.29.2.5 Node() [5/5] 1	67
7.29.2.6 ~Node()	68
7.29.3 Member Function Documentation	68
7.29.3.1 get_parent()	68
7.29.3.2 is_leaf()	68
7.29.3.3 noffspring()	68
7.29.4 Member Data Documentation	68
7.29.4.1 annotations	68
7.29.4.2 array	69
7.29.4.3 arrays	69
7.29.4.4 duplication	69
7.29.4.5 id	69
7.29.4.6 narray	69
7.29.4.7 offspring	70
7.29.4.8 ord	70
7.29.4.9 parent	70
7.29.4.10 probability	70

7.29.4.11 subtree_prob	70
7.29.4.12 visited	71
7.30 NodeData Class Reference	71
7.30.1 Detailed Description	71
7.30.2 Constructor & Destructor Documentation	71
7.30.2.1 NodeData()	71
7.30.3 Member Data Documentation	72
7.30.3.1 blengths	72
7.30.3.2 duplication	72
7.30.3.3 states	72
7.31 PhyloCounterData Class Reference	72
7.31.1 Detailed Description	73
7.31.2 Constructor & Destructor Documentation	73
7.31.2.1 PhyloCounterData() [1/2]	73
7.31.2.2 PhyloCounterData() [2/2]	73
7.31.3 Member Function Documentation	73
7.31.3.1 at()	73
7.31.3.2 begin()	73
7.31.3.3 empty()	74
7.31.3.4 end()	74
7.31.3.5 get_counters()	74
7.31.3.6 operator()()	74
7.31.3.7 operator[]()	74
7.31.3.8 push_back()	74
7.31.3.9 reserve()	75
7.31.3.10 shrink_to_fit()	75
7.31.3.11 size()	75
7.32 PhyloRuleDynData Class Reference	75
7.32.1 Detailed Description	75
7.32.2 Constructor & Destructor Documentation	76
7.32.2.1 PhyloRuleDynData()	76
$7.32.2.2 \sim$ PhyloRuleDynData()	76
7.32.3 Member Data Documentation	76
7.32.3.1 counts	76
7.32.3.2 duplication	76
7.32.3.3 lb	76
7.32.3.4 pos	77
7.32.3.5 ub	77
7.33 PowerSet< Array_Type, Data_Rule_Type > Class Template Reference	77
7.33.1 Detailed Description	78
7.33.2 Constructor & Destructor Documentation	79
7.33.2.1 PowerSet() [1/3]	79

7.33.2.2 PowerSet() [2/3]	179
7.33.2.3 PowerSet() [3/3]	179
7.33.2.4 ~PowerSet()	179
7.33.3 Member Function Documentation	179
7.33.3.1 add_rule() [1/2]	180
7.33.3.2 add_rule() [2/2]	180
7.33.3.3 begin()	180
7.33.3.4 calc()	180
7.33.3.5 end()	180
7.33.3.6 get_data()	181
7.33.3.7 get_data_ptr()	181
7.33.3.8 init_support()	181
7.33.3.9 operator[]()	181
7.33.3.10 reset()	181
7.33.3.11 size()	182
7.33.4 Member Data Documentation	182
7.33.4.1 coordinates_free	182
7.33.4.2 coordinates_locked	182
7.33.4.3 data	182
7.33.4.4 EmptyArray	182
7.33.4.5 M	183
7.33.4.6 N	183
7.33.4.7 n_free	183
7.33.4.8 n_locked	183
7.33.4.9 rules	183
7.33.4.10 rules_deleted	184
7.34 Progress Class Reference	184
7.34.1 Detailed Description	184
7.34.2 Constructor & Destructor Documentation	184
7.34.2.1 Progress()	184
7.34.2.2 ~Progress()	185
7.34.3 Member Function Documentation	185
7.34.3.1 end()	185
7.34.3.2 next()	185
7.35 Rule < Array_Type, Data_Type > Class Template Reference	185
7.35.1 Detailed Description	186
7.35.2 Constructor & Destructor Documentation	186
7.35.2.1 Rule() [1/2]	186
7.35.2.2 Rule() [2/2]	186
7.35.2.3 ~Rule()	187
7.35.3 Member Function Documentation	187
7.35.3.1 D()	187

7.35.3.2 operator()()	187
7.36 Rules < Array_Type, Data_Type > Class Template Reference	187
7.36.1 Detailed Description	188
7.36.2 Constructor & Destructor Documentation	188
7.36.2.1 Rules() [1/2] 1	188
7.36.2.2 Rules() [2/2] 1	188
7.36.2.3 ~Rules()	189
7.36.3 Member Function Documentation	189
7.36.3.1 add_rule() [1/2]1	189
7.36.3.2 add_rule() [2/2]1	189
7.36.3.3 get_seq()	189
7.36.3.4 operator()()	190
7.36.3.5 operator=()	190
7.36.3.6 size()	190
7.37 StatsCounter< Array_Type, Data_Type > Class Template Reference	191
7.37.1 Detailed Description	191
7.37.2 Constructor & Destructor Documentation	191
7.37.2.1 StatsCounter() [1/3]	191
7.37.2.2 StatsCounter() [2/3]	192
7.37.2.3 StatsCounter() [3/3]	192
7.37.2.4 ~StatsCounter()	192
7.37.3 Member Function Documentation	192
7.37.3.1 add_counter()	192
7.37.3.2 count_all()	193
7.37.3.3 count_current()	193
7.37.3.4 count_init()	193
7.37.3.5 get_counters()	193
7.37.3.6 get_descriptions()	193
7.37.3.7 get_names()	193
7.37.3.8 reset_array()	193
7.37.3.9 set_counters()	194
7.37.3.10 size()	194
7.38 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class Template Reference	194
7.38.1 Detailed Description	196
7.38.2 Constructor & Destructor Documentation	196
7.38.2.1 Support() [1/3]	196
7.38.2.2 Support() [2/3]	197
7.38.2.3 Support() [3/3]	197
7.38.2.4 ~Support()	197
7.38.3 Member Function Documentation	197
7.38.3.1 add_counter()	197

7.38.3.2 add_rule() [1/2] 19	8
7.38.3.3 add_rule() [2/2]19	8
7.38.3.4 add_rule_dyn() [1/2]	8
7.38.3.5 add_rule_dyn() [2/2]	8
7.38.3.6 calc()	8
7.38.3.7 eval_rules_dyn()	9
7.38.3.8 get_counters()	9
7.38.3.9 get_counts()	9
7.38.3.10 get_current_stats()	9
7.38.3.11 get_data()	0
7.38.3.12 get_rules()	0
7.38.3.13 get_rules_dyn()	0
7.38.3.14 init_support()	0
7.38.3.15 print()	0
7.38.3.16 reset_array() [1/2]	1
7.38.3.17 reset_array() [2/2]	1
7.38.3.18 set_counters()	1
7.38.3.19 set_rules()	1
7.38.3.20 set_rules_dyn()	1
7.38.4 Member Data Documentation	1
7.38.4.1 change_stats	2
7.38.4.2 coordiantes_n_free	2
7.38.4.3 coordiantes_n_locked	2
7.38.4.4 coordinates_free	2
7.38.4.5 coordinates_locked	
7.38.4.6 current_stats	
7.38.4.7 delete_counters	3
7.38.4.8 delete_rules	3
7.38.4.9 delete_rules_dyn	3
7.38.4.10 hashes	3
7.38.4.11 hashes_initialized	4
7.38.4.12 M	4
7.38.4.13 max_num_elements	
7.38.4.14 N	
7.38.4.15 n_counters	4
7.39 vecHasher < T > Struct Template Reference	5
7.39.1 Detailed Description	5
7.39.2 Member Function Documentation	5
7.39.2.1 operator()()	5
8 File Documentation 20	7
8.1 include/barry/barray-bones.hpp File Reference	7

8.1.1 Macro Definition Documentation	208
8.1.1.1 BARRAY_BONES_HPP	208
8.2 include/barry/barray-iterator.hpp File Reference	208
8.3 include/barry/barray-meat-operators.hpp File Reference	209
8.3.1 Macro Definition Documentation	210
8.3.1.1 BARRAY_TEMPLATE	210
8.3.1.2 BARRAY_TEMPLATE_ARGS	210
8.3.1.3 BARRAY_TYPE	210
8.3.1.4 BARRY_BARRAY_MEAT_OPERATORS_HPP	211
8.3.1.5 COL	211
8.3.1.6 ROW	211
8.3.2 Function Documentation	211
8.3.2.1 BARRAY_TEMPLATE() [1/6]	211
8.3.2.2 BARRAY_TEMPLATE() [2/6]	211
8.3.2.3 BARRAY_TEMPLATE() [3/6]	212
8.3.2.4 BARRAY_TEMPLATE() [4/6]	212
8.3.2.5 BARRAY_TEMPLATE() [5/6]	212
8.3.2.6 BARRAY_TEMPLATE() [6/6]	212
8.3.2.7 BARRAY_TEMPLATE_ARGS()	212
8.3.2.8 BARRAY_TYPE()	212
8.3.2.9 for()	213
8.3.2.10 operator()()	213
8.3.3 Variable Documentation	213
8.3.3.1 rhs	213
8.3.3.2 this	213
8.4 include/barry/barray-meat.hpp File Reference	214
8.4.1 Macro Definition Documentation	216
8.4.1.1 BARRAY_TEMPLATE	216
8.4.1.2 BARRAY_TEMPLATE_ARGS	216
8.4.1.3 BARRAY_TYPE	217
8.4.1.4 COL	217
8.4.1.5 ROW	217
8.4.2 Function Documentation	217
8.4.2.1 ans()	217
8.4.2.2 BARRAY_TEMPLATE() [1/24]	217
8.4.2.3 BARRAY_TEMPLATE() [2/24]	218
8.4.2.4 BARRAY_TEMPLATE() [3/24]	218
8.4.2.5 BARRAY_TEMPLATE() [4/24]	218
8.4.2.6 BARRAY_TEMPLATE() [5/24]	
8.4.2.7 BARRAY_TEMPLATE() [6/24]	218
8.4.2.8 BARRAY_TEMPLATE() [7/24]	218
8.4.2.9 BARRAY_TEMPLATE() [8/24]	219

8.4.2.10 BARRAY_TEMPLATE() [9/24]
8.4.2.11 BARRAY_TEMPLATE() [10/24]
8.4.2.12 BARRAY_TEMPLATE() [11/24]
8.4.2.13 BARRAY_TEMPLATE() [12/24]
8.4.2.14 BARRAY_TEMPLATE() [13/24]
8.4.2.15 BARRAY_TEMPLATE() [14/24]
8.4.2.16 BARRAY_TEMPLATE() [15/24]
8.4.2.17 BARRAY_TEMPLATE() [16/24]
8.4.2.18 BARRAY_TEMPLATE() [17/24]
8.4.2.19 BARRAY_TEMPLATE() [18/24]
8.4.2.20 BARRAY_TEMPLATE() [19/24]
8.4.2.21 BARRAY_TEMPLATE() [20/24]
8.4.2.22 BARRAY_TEMPLATE() [21/24]
8.4.2.23 BARRAY_TEMPLATE() [22/24]
8.4.2.24 BARRAY_TEMPLATE() [23/24]
8.4.2.25 BARRAY_TEMPLATE() [24/24]
8.4.2.26 COL()
8.4.2.27 for() [1/3]
8.4.2.28 for() [2/3]
8.4.2.29 for() [3/3]
8.4.2.30 if() [1/17]
8.4.2.31 if() [2/17]
8.4.2.32 if() [3/17]
8.4.2.33 if() [4/17]
8.4.2.34 if() [5/17]
8.4.2.35 if() [6/17]
8.4.2.36 if() [7/17]
8.4.2.37 if() [8/17]
8.4.2.38 if() [9/17]
8.4.2.39 if() [10/17]
8.4.2.40 if() [11/17]
8.4.2.41 if() [12/17]
8.4.2.42 if() [13/17]
8.4.2.43 if() [14/17]
8.4.2.44 if() [15/17]
8.4.2.45 if() [16/17]
8.4.2.46 if() [17/17]
8.4.2.47 M()
8.4.2.48 resize() [1/2]
8.4.2.49 resize() [2/2]
8.4.2.50 return()
8.4.2.51 ROW() [1/2]

8.4.2.52 ROW() [2/2]	26
8.4.3 Variable Documentation	26
8.4.3.1 add	26
8.4.3.2 ans	26
8.4.3.3 Array	26
8.4.3.4 check_bounds	27
8.4.3.5 check_exists	27
8.4.3.6 col0	27
8.4.3.7 const	27
8.4.3.8 copy_data	28
8.4.3.9 data	28
8.4.3.10 delete_data	28
8.4.3.11 delete_data	28
8.4.3.12 else	28
8.4.3.13 false	29
8.4.3.14 first	29
8.4.3.15 i1	29
8.4.3.16 j	29
8.4.3.17 j0	29
8.4.3.18 j1	29
8.4.3.19 M	30
8.4.3.20 M	30
8.4.3.21 N	30
8.4.3.22 NCells	30
8.4.3.23 report	30
8.4.3.24 return	31
8.4.3.25 row0	31
8.4.3.26 search	31
8.4.3.27 source	31
8.4.3.28 target	31
8.4.3.29 v	31
8.4.3.30 value	32
8.5 include/barry/barraycell-bones.hpp File Reference	32
8.6 include/barry/barraycell-meat.hpp File Reference	33
8.7 include/barry/barraydense-bones.hpp File Reference	33
8.7.1 Macro Definition Documentation	35
8.7.1.1 BARRY_BARRAYDENSE_BONES_HPP	35
8.8 include/barry/barraydense-meat-operators.hpp File Reference	35
8.8.1 Macro Definition Documentation	36
8.8.1.1 BARRY_BARRAYDENSE_MEAT_OPERATORS_HPP	36
8.8.1.2 BDENSE_TEMPLATE	36
8.8.1.3 BDENSE_TEMPLATE_ARGS	36

8.8.1.4 BDENSE_TYPE	37
8.8.1.5 COL	37
8.8.1.6 POS	37
8.8.1.7 POS_N	37
8.8.1.8 ROW	37
8.8.2 Function Documentation	37
8.8.2.1 BDENSE_TEMPLATE() [1/4]	38
8.8.2.2 BDENSE_TEMPLATE() [2/4]	38
8.8.2.3 BDENSE_TEMPLATE() [3/4]	38
8.8.2.4 BDENSE_TEMPLATE() [4/4]	
8.8.2.5 BDENSE_TEMPLATE_ARGS()	38
8.8.2.6 BDENSE_TYPE()	38
8.9 include/barry/barraydense-meat.hpp File Reference	39
8.9.1 Macro Definition Documentation	
8.9.1.1 BDENSE_TEMPLATE	41
8.9.1.2 BDENSE_TEMPLATE_ARGS	42
8.9.1.3 BDENSE_TYPE	42
8.9.1.4 COL	42
8.9.1.5 POS	42
8.9.1.6 POS_N	42
8.9.1.7 ROW	43
8.9.1.8 ZERO_CELL	43
8.9.2 Function Documentation	43
8.9.2.1 ans()	43
8.9.2.2 BDENSE_TEMPLATE() [1/39]	43
8.9.2.3 BDENSE_TEMPLATE() [2/39]	43
8.9.2.4 BDENSE_TEMPLATE() [3/39]	43
8.9.2.5 BDENSE_TEMPLATE() [4/39]	44
8.9.2.6 BDENSE_TEMPLATE() [5/39]	44
8.9.2.7 BDENSE_TEMPLATE() [6/39]	44
8.9.2.8 BDENSE_TEMPLATE() [7/39]	44
8.9.2.9 BDENSE_TEMPLATE() [8/39]	44
8.9.2.10 BDENSE_TEMPLATE() [9/39]	45
8.9.2.11 BDENSE_TEMPLATE() [10/39]	45
8.9.2.12 BDENSE_TEMPLATE() [11/39]	45
8.9.2.13 BDENSE_TEMPLATE() [12/39]	45
8.9.2.14 BDENSE_TEMPLATE() [13/39]	45
8.9.2.15 BDENSE_TEMPLATE() [14/39]	46
8.9.2.16 BDENSE_TEMPLATE() [15/39]	46
8.9.2.17 BDENSE_TEMPLATE() [16/39]	46
8.9.2.18 BDENSE_TEMPLATE() [17/39]	46
8.9.2.19 BDENSE_TEMPI ATE() [18/39]	46

8.9.2.20 BDENSE_TEMPLATE() [19/	/39]	 	 	247
8.9.2.21 BDENSE_TEMPLATE() [20/	/39]	 	 	247
8.9.2.22 BDENSE_TEMPLATE() [21/	/39]	 	 	247
8.9.2.23 BDENSE_TEMPLATE() [22/	/39]	 	 	247
8.9.2.24 BDENSE_TEMPLATE() [23/	/39]	 	 	247
8.9.2.25 BDENSE_TEMPLATE() [24/	/39]	 	 	248
8.9.2.26 BDENSE_TEMPLATE() [25/	/39]	 	 	248
8.9.2.27 BDENSE_TEMPLATE() [26/	/39]	 	 	248
8.9.2.28 BDENSE_TEMPLATE() [27/				
8.9.2.29 BDENSE_TEMPLATE() [28/				
8.9.2.30 BDENSE_TEMPLATE() [29/	/39]	 	 	249
8.9.2.31 BDENSE_TEMPLATE() [30/				
8.9.2.32 BDENSE_TEMPLATE() [31/				
8.9.2.33 BDENSE_TEMPLATE() [32/				
8.9.2.34 BDENSE_TEMPLATE() [33/				
8.9.2.35 BDENSE_TEMPLATE() [34/				
8.9.2.36 BDENSE_TEMPLATE() [35/				
8.9.2.37 BDENSE_TEMPLATE() [36/				
8.9.2.38 BDENSE_TEMPLATE() [37/				
8.9.2.39 BDENSE_TEMPLATE() [38/				
8.9.2.40 BDENSE_TEMPLATE() [39/				
8.9.2.41 for()				
8.9.2.42 if() [1/4]				
8.9.2.43 if() [2/4]				
8.9.2.44 if() [3/4]				
8.9.2.45 if() [4/4]				
8.9.2.46 insert_cell() [1/2]				
8.9.2.47 insert_cell() [2/2]				
8.9.2.48 M()				
8.9.2.49 resize() [1/6]				
8.9.2.50 resize() [2/6]				
8.9.2.51 resize() [3/6]				
8.9.2.52 resize() [4/6]				
8.9.2.53 resize() [5/6]				
8.9.2.54 resize() [6/6]				
8.9.2.55 rm_cell() [1/3]				
8.9.2.56 rm_cell() [2/3]				
8.9.2.57 rm_cell() [3/3]				
8.9.2.58 va_end()				
8.9.2.59 va_start()				
8.9.2.60 vprintf()				
8.9.3 Variable Documentation		 	 	254

8.9.3.1 add	54
8.9.3.2 ans	54
8.9.3.3 check_bounds	54
8.9.3.4 check_exists	55
8.9.3.5 col	55
8.9.3.6 const	55
8.9.3.7 copy_data	55
8.9.3.8 data	55
8.9.3.9 delete_data	56
8.9.3.10 delete_data	56
8.9.3.11 el	56
8.9.3.12 el_colsums	56
8.9.3.13 el_rowsums	56
8.9.3.14 else	57
8.9.3.15 false	57
8.9.3.16 i1	57
8.9.3.17 j	57
8.9.3.18 j0	57
8.9.3.19 j1	57
8.9.3.20 M	58
8.9.3.21 M	58
8.9.3.22 N	58
8.9.3.23 report	58
8.9.3.24 return	58
8.9.3.25 source	59
8.9.3.26 target	59
8.9.3.27 v	59
8.9.3.28 val0	59
8.9.3.29 val1	59
8.9.3.30 value	59
8.10 include/barry/barraydensecell-bones.hpp File Reference	60
8.10.1 Macro Definition Documentation	60
8.10.1.1 POS	61
8.11 include/barry/barraydensecell-meat.hpp File Reference	61
8.11.1 Macro Definition Documentation	62
8.11.1.1 POS	62
8.12 include/barry/barraydensecol-bones.hpp File Reference	62
8.12.1 Macro Definition Documentation	63
8.12.1.1 POS	63
8.12.1.2 POS_N	63
8.12.1.3 ZERO_CELL	63
8.13 include/barry/barraydenserow-bones.hpp File Reference	63

8.13.1 Macro Definition Documentation	34
8.13.1.1 POS	34
8.13.1.2 POS_N	34
8.13.1.3 ZERO_CELL	34
8.14 include/barry/barrayrow-bones.hpp File Reference	35
8.15 include/barry/barrayrow-meat.hpp File Reference	35
8.15.1 Macro Definition Documentation	36
8.15.1.1 BARRY_BARRAYROW_MEAT_HPP	36
8.15.1.2 BROW_TEMPLATE	37
8.15.1.3 BROW_TEMPLATE_ARGS	37
8.15.1.4 BROW_TYPE	37
8.15.2 Function Documentation	37
8.15.2.1 BROW_TEMPLATE() [1/5]	37
8.15.2.2 BROW_TEMPLATE() [2/5]	37
8.15.2.3 BROW_TEMPLATE() [3/5]	38
8.15.2.4 BROW_TEMPLATE() [4/5]	38
8.15.2.5 BROW_TEMPLATE() [5/5]	38
8.16 include/barry/barrayvector-bones.hpp File Reference	38
8.17 include/barry/barrayvector-meat.hpp File Reference	39
8.17.1 Macro Definition Documentation	70
8.17.1.1 BARRY_BARRAYVECTOR_MEAT_HPP	70
8.18 include/barry/barry-configuration.hpp File Reference	70
8.18.1 Macro Definition Documentation	71
8.18.1.1 BARRY_CHECK_SUPPORT	71
8.18.1.2 BARRY_ISFINITE	71
8.18.1.3 BARRY_MAX_NUM_ELEMENTS	71
8.18.1.4 BARRY_SAFE_EXP	71
8.18.1.5 printf_barry	71
8.18.2 Typedef Documentation	71
8.18.2.1 Map	72
8.19 include/barry/barry-debug.hpp File Reference	72
8.19.1 Macro Definition Documentation	72
8.19.1.1 BARRY_DEBUG_LEVEL	72
8.20 include/barry/barry-macros.hpp File Reference	72
0.00 () 1	73
8.20.1 Macro Definition Documentation	
8.20.1 Macro Definition Documentation	
	73
8.20.1.1 BARRY_ONE	73 73
8.20.1.1 BARRY_ONE	73 73 73
8.20.1.1 BARRY_ONE 27 8.20.1.2 BARRY_ONE_DENSE 27 8.20.1.3 BARRY_UNUSED 27	73 73 73 73
8.20.1.1 BARRY_ONE 27 8.20.1.2 BARRY_ONE_DENSE 27 8.20.1.3 BARRY_UNUSED 27 8.20.1.4 BARRY_ZERO 27	73 73 73 73 74

8.21.1.1 BARRY_HPP
8.21.1.2 BARRY_VERSION
8.21.1.3 COUNTER_FUNCTION
8.21.1.4 COUNTER_LAMBDA
8.21.1.5 RULE_FUNCTION
8.21.1.6 RULE_LAMBDA
8.22 include/barry/cell-bones.hpp File Reference
8.23 include/barry/cell-meat.hpp File Reference
8.24 include/barry/col-bones.hpp File Reference
8.25 include/barry/counters-bones.hpp File Reference
8.26 include/barry/counters-meat.hpp File Reference
8.26.1 Macro Definition Documentation
8.26.1.1 COUNTER_TEMPLATE
8.26.1.2 COUNTER_TEMPLATE_ARGS
8.26.1.3 COUNTER_TYPE
8.26.1.4 COUNTERS_TEMPLATE
8.26.1.5 COUNTERS_TEMPLATE_ARGS
8.26.1.6 COUNTERS_TYPE
8.26.2 Function Documentation
8.26.2.1 count_fun()
8.26.2.2 COUNTER_TEMPLATE() [1/7]
8.26.2.3 COUNTER_TEMPLATE() [2/7]
8.26.2.4 COUNTER_TEMPLATE() [3/7]
8.26.2.5 COUNTER_TEMPLATE() [4/7]
8.26.2.6 COUNTER_TEMPLATE() [5/7]
8.26.2.7 COUNTER_TEMPLATE() [6/7]
8.26.2.8 COUNTER_TEMPLATE() [7/7]
8.26.2.9 Counters()
8.26.2.10 COUNTERS_TEMPLATE() [1/7]
8.26.2.11 COUNTERS_TEMPLATE() [2/7]
8.26.2.12 COUNTERS_TEMPLATE() [3/7]
8.26.2.13 COUNTERS_TEMPLATE() [4/7]
8.26.2.14 COUNTERS_TEMPLATE() [5/7]
8.26.2.15 COUNTERS_TEMPLATE() [6/7]
8.26.2.16 COUNTERS_TEMPLATE() [7/7]
8.26.2.17 data()
8.26.2.18 desc()
8.26.2.19 init_fun() [1/3]
8.26.2.20 init_fun() [2/3]
8.26.2.21 init_fun() [3/3]
8.26.2.22 name()
8.26.3 Variable Documentation 287

	8.26.3.1 count_fun	 		287
	8.26.3.2 counter	 		287
	8.26.3.3 counter	 		287
	8.26.3.4 data	 		288
	8.26.3.5 desc	 		288
	8.26.3.6 i	 		288
	8.26.3.7 init_fun	 		288
	8.26.3.8 j	 		288
	8.26.3.9 name	 		289
	8.26.3.10 noexcept	 		289
	8.26.3.11 return	 		289
	8.26.3.12 this	 		289
8.27 include/b	parry/counters/defm.hpp File Reference	 		289
8.27.1 N	Macro Definition Documentation	 		291
	8.27.1.1 DEFM_COUNTER	 		291
	8.27.1.2 DEFM_COUNTER_LAMBDA	 		291
	8.27.1.3 DEFM_RULE	 		291
	8.27.1.4 DEFM_RULE_LAMBDA	 		292
8.27.2 T	ypedef Documentation	 		292
	8.27.2.1 DEFMArray	 		292
	8.27.2.2 DEFMCounter	 		292
	8.27.2.3 DEFMCounters	 		292
	8.27.2.4 DEFMModel	 		292
	8.27.2.5 DEFMRule	 		293
	8.27.2.6 DEFMRules	 		293
	8.27.2.7 DEFMStatsCounter	 		293
	8.27.2.8 DEFMSupport	 		293
8.28 include/b	parry/models/defm.hpp File Reference	 		293
8.29 include/b	parry/counters/network-css.hpp File Reference	 		294
8.29.1 N	Macro Definition Documentation	 		295
	8.29.1.1 CSS_APPEND	 		295
	8.29.1.2 CSS_CASE_ELSE	 		296
	8.29.1.3 CSS_CASE_PERCEIVED	 		296
	8.29.1.4 CSS_CASE_TRUTH	 		296
	8.29.1.5 CSS_CHECK_SIZE	 		296
	8.29.1.6 CSS_CHECK_SIZE_INIT	 		296
	8.29.1.7 CSS_NET_COUNTER_LAMBDA_INIT	 		297
	8.29.1.8 CSS_PERCEIVED_CELLS	 		297
	8.29.1.9 CSS_SIZE	 		297
	8.29.1.10 CSS_TRUE_CELLS	 		297
8.29.2 F	Function Documentation	 		297
	8.29.2.1 counter css census01()	 		298

8.29.2.2 counter_css_census02()	298
8.29.2.3 counter_css_census03()	298
8.29.2.4 counter_css_census04()	298
8.29.2.5 counter_css_census05()	299
8.29.2.6 counter_css_census06()	299
8.29.2.7 counter_css_census07()	299
8.29.2.8 counter_css_census08()	299
8.29.2.9 counter_css_census09()	300
8.29.2.10 counter_css_census10()	300
8.29.2.11 counter_css_completely_false_recip_comiss()	300
8.29.2.12 counter_css_completely_false_recip_omiss()	300
8.29.2.13 counter_css_mixed_recip()	301
8.29.2.14 counter_css_partially_false_recip_commi()	301
8.29.2.15 counter_css_partially_false_recip_omiss()	301
8.30 include/barry/counters/network.hpp File Reference	302
8.30.1 Macro Definition Documentation	304
8.30.1.1 BARRY_ZERO_NETWORK	305
8.30.1.2 BARRY_ZERO_NETWORK_DENSE	305
8.30.1.3 NET_C_DATA_IDX	305
8.30.1.4 NET_C_DATA_NUM	305
8.30.1.5 NETWORK_COUNTER	305
8.30.1.6 NETWORK_COUNTER_LAMBDA	306
8.30.1.7 NETWORK_RULE	306
8.30.1.8 NETWORK_RULE_LAMBDA	306
8.30.1.9 NETWORKDENSE_COUNTER_LAMBDA	306
8.30.2 Typedef Documentation	307
8.30.2.1 NetCounter	307
8.30.2.2 NetCounters	307
8.30.2.3 NetModel	307
8.30.2.4 NetRule	307
8.30.2.5 NetRules	307
8.30.2.6 NetStatsCounter	308
8.30.2.7 NetSupport	308
8.30.2.8 Network	308
8.30.2.9 NetworkDense	308
8.30.3 Function Documentation	308
8.30.3.1 rules_zerodiag()	308
8.31 include/barry/counters/phylo.hpp File Reference	309
8.31.1 Macro Definition Documentation	311
8.31.1.1 DEFAULT_DUPLICATION	311
8.31.1.2 DUPL_DUPL	311
8 31 1 3 DUDLETH	211

8.31.1.4 DUPL_SPEC	2
8.31.1.5 IF_MATCHES	2
8.31.1.6 IF_NOTMATCHES	2
8.31.1.7 IS_DUPLICATION	2
8.31.1.8 IS_EITHER	2
8.31.1.9 IS_SPECIATION	3
8.31.1.10 MAKE_DUPL_VARS	3
8.31.1.11 PHYLO_CHECK_MISSING	3
8.31.1.12 PHYLO_COUNTER_LAMBDA	3
8.31.1.13 PHYLO_RULE_DYN_LAMBDA	4
8.31.2 Typedef Documentation	4
8.31.2.1 PhyloArray	4
8.31.2.2 PhyloCounter	4
8.31.2.3 PhyloCounters	4
8.31.2.4 PhyloModel	4
8.31.2.5 PhyloPowerSet	5
8.31.2.6 PhyloRule	5
8.31.2.7 PhyloRuleData	5
8.31.2.8 PhyloRuleDyn	5
8.31.2.9 PhyloRules	5
8.31.2.10 PhyloRulesDyn	5
8.31.2.11 PhyloStatsCounter	6
8.31.2.12 PhyloSupport	6
8.31.3 Function Documentation	6
8.31.3.1 get_last_name()	6
8.32 include/barry/freqtable.hpp File Reference	6
8.33 include/barry/model-bones.hpp File Reference	7
8.33.1 Function Documentation	9
8.33.1.1 keygen_default()	9
8.34 include/barry/model-meat.hpp File Reference	9
8.34.1 Macro Definition Documentation	20
8.34.1.1 MODEL_TEMPLATE	20
8.34.1.2 MODEL_TEMPLATE_ARGS	20
8.34.1.3 MODEL_TYPE	20
8.34.2 Function Documentation	20
8.34.2.1 likelihood_()	20
8.34.2.2 MODEL_TEMPLATE() [1/2]	21
8.34.2.3 MODEL_TEMPLATE() [2/2]	21
8.34.2.4 update_normalizing_constant()	21
8.35 include/barry/models/defm/defm-bones.hpp File Reference	21
8.35.1 Macro Definition Documentation	22
8.35.1.1 INITIALIZED	2

8.35.2 Function Documentation
8.35.2.1 keygen_full()
8.35.2.2 RULE_FUNCTION()
8.35.2.3 vec_diff()
8.36 include/barry/models/defm/defm-meat.hpp File Reference
8.37 include/barry/models/geese.hpp File Reference
8.38 include/barry/models/geese/flock-bones.hpp File Reference
8.39 include/barry/models/geese/flock-meat.hpp File Reference
8.40 include/barry/models/geese/geese-bones.hpp File Reference
8.40.1 Macro Definition Documentation
8.40.1.1 INITIALIZED
8.40.2 Function Documentation
8.40.2.1 keygen_full()
8.40.2.2 RULE_FUNCTION()
8.40.2.3 vec_diff()
8.40.2.4 vector_caster()
8.41 include/barry/models/geese/geese-meat-constructors.hpp File Reference
8.42 include/barry/models/geese/geese-meat-likelihood.hpp File Reference
8.43 include/barry/models/geese/geese-meat-likelihood_exhaust.hpp File Reference
8.44 include/barry/models/geese/geese-meat-predict.hpp File Reference
8.45 include/barry/models/geese/geese-meat-predict_exhaust.hpp File Reference
8.46 include/barry/models/geese/geese-meat-predict_sim.hpp File Reference
8.47 include/barry/models/geese/geese-meat-simulate.hpp File Reference
8.48 include/barry/models/geese/geese-meat.hpp File Reference
8.49 include/barry/models/geese/geese-node-bones.hpp File Reference
8.50 include/barry/powerset-bones.hpp File Reference
8.51 include/barry/powerset-meat.hpp File Reference
8.52 include/barry/progress.hpp File Reference
8.52.1 Macro Definition Documentation
8.52.1.1 BARRY_PROGRESS_BAR_WIDTH
8.53 include/barry/rules-bones.hpp File Reference
8.53.1 Function Documentation
8.53.1.1 rule_fun_default()
8.54 include/barry/rules-meat.hpp File Reference
8.55 include/barry/statscounter-bones.hpp File Reference
8.56 include/barry/statscounter-meat.hpp File Reference
8.56.1 Macro Definition Documentation
8.56.1.1 STATSCOUNTER_TEMPLATE
8.56.1.2 STATSCOUNTER_TEMPLATE_ARGS
8.56.1.3 STATSCOUNTER_TYPE
8.56.2 Function Documentation
8.56.2.1 clear()

8.56.2.2 for()	. 340
8.56.2.3 resize()	. 340
8.56.2.4 STATSCOUNTER_TEMPLATE() [1/9]	. 340
8.56.2.5 STATSCOUNTER_TEMPLATE() [2/9]	. 340
8.56.2.6 STATSCOUNTER_TEMPLATE() [3/9]	. 341
8.56.2.7 STATSCOUNTER_TEMPLATE() [4/9]	. 341
8.56.2.8 STATSCOUNTER_TEMPLATE() [5/9]	. 341
8.56.2.9 STATSCOUNTER_TEMPLATE() [6/9]	. 341
8.56.2.10 STATSCOUNTER_TEMPLATE() [7/9]	. 341
8.56.2.11 STATSCOUNTER_TEMPLATE() [8/9]	. 341
8.56.2.12 STATSCOUNTER_TEMPLATE() [9/9]	. 342
8.56.3 Variable Documentation	. 342
8.56.3.1 counter	. 342
8.56.3.2 counter_deleted	. 342
8.56.3.3 counters	. 342
8.56.3.4 counters	. 342
8.56.3.5 current_stats	. 343
8.56.3.6 EmptyArray	. 343
8.56.3.7 f	. 343
8.56.3.8 j	. 343
8.56.3.9 return	. 343
8.57 include/barry/support-bones.hpp File Reference	. 344
8.58 include/barry/support-meat.hpp File Reference	. 345
8.58.1 Macro Definition Documentation	. 347
8.58.1.1 BARRY_SUPPORT_MEAT_HPP	. 347
8.58.1.2 SUPPORT_TEMPLATE	. 347
8.58.1.3 SUPPORT_TEMPLATE_ARGS	. 347
8.58.1.4 SUPPORT_TYPE	. 347
8.58.2 Function Documentation	. 347
8.58.2.1 calc_backend_dense()	. 348
8.58.2.2 calc_backend_sparse()	. 348
8.58.2.3 for()	. 348
8.58.2.4 if() [1/3]	. 348
8.58.2.5 if() [2/3]	. 348
8.58.2.6 if() [3/3]	. 348
8.58.2.7 insert_cell() [1/2]	. 349
8.58.2.8 insert_cell() [2/2]	. 349
8.58.2.9 rm_cell()	. 349
8.58.2.10 SUPPORT_TEMPLATE() [1/17]	. 349
8.58.2.11 SUPPORT_TEMPLATE() [2/17]	. 349
8.58.2.12 SUPPORT_TEMPLATE() [3/17]	. 350
8.58.2.13 SUPPORT_TEMPLATE() [4/17]	. 350

8.58.2.14 SUPPORT_TEMPLATE() [5/17]	 350
8.58.2.15 SUPPORT_TEMPLATE() [6/17]	 350
8.58.2.16 SUPPORT_TEMPLATE() [7/17]	 350
8.58.2.17 SUPPORT_TEMPLATE() [8/17]	 350
8.58.2.18 SUPPORT_TEMPLATE() [9/17]	 351
8.58.2.19 SUPPORT_TEMPLATE() [10/17]	 351
8.58.2.20 SUPPORT_TEMPLATE() [11/17]	 351
8.58.2.21 SUPPORT_TEMPLATE() [12/17]	 351
8.58.2.22 SUPPORT_TEMPLATE() [13/17]	 351
8.58.2.23 SUPPORT_TEMPLATE() [14/17]	 352
8.58.2.24 SUPPORT_TEMPLATE() [15/17]	 352
8.58.2.25 SUPPORT_TEMPLATE() [16/17]	 352
8.58.2.26 SUPPORT_TEMPLATE() [17/17]	 352
8.58.3 Variable Documentation	 352
8.58.3.1 array_bank	 352
8.58.3.2 change_stats_different	 353
8.58.3.3 coord_i	 353
8.58.3.4 coord_j	 353
8.58.3.5 counters	
8.58.3.6 counters	 353
8.58.3.7 delete_counters	 353
8.58.3.8 delete_rules	 354
8.58.3.9 delete_rules_dyn	 354
8.58.3.10 else	 354
8.58.3.11 f	 354
8.58.3.12 hashes	 354
8.58.3.13 return	 355
8.58.3.14 rules	 355
8.58.3.15 rules	
8.58.3.16 rules_dyn	 355
8.58.3.17 stats_bank	
8.58.3.18 tmp_chng	 356
8.59 include/barry/typedefs.hpp File Reference	
8.59.1 Typedef Documentation	
8.59.1.1 Col_type	 358
8.59.1.2 Counter_fun_type	 358
8.59.1.3 Counts_type	
8.59.1.4 MapVec_type	 358
8.59.1.5 Row_type	
8.59.1.6 Rule_fun_type	 359
8.59.1.7 uint	 359
8.59.2 Function Documentation	 359

8.59.2.1 vec_equal()	359
8.59.2.2 vec_equal_approx()	360
8.59.2.3 vec_inner_prod() [1/2]	360
8.59.2.4 vec_inner_prod() [2/2]	360

xxxvii

360

361

Index

Chapter 1

Main Page

Barry: your to-go motif accountant

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

Among the key features included in barry, we have:

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

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

Examples

Counting statistics in a graph

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

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

2 Main Page

```
// Creating network of size six with five ties
  netcounters::Network net(
       6, 6,
      {0, 0, 4, 4, 2, 0, 1},
      {1, 2, 0, 2, 4, 0, 1}
  // How does this looks like?
  net.print("Current view");
  // Adding extra ties
  net += {1, 0};
net(2, 0) = true;
  // And removing a couple
  net(0, 0) = false;
net -= {1, 1};
  net.print("New view");
  // Initializing the data. The program deals with freing the memory
  net.set_data(new netcounters::NetworkData, true);
  // Creating counter object for the network and adding stats to count
  netcounters::NetStatsCounter counter(&net);
  netcounters::counter_edges(counter.counters);
  netcounters::counter_ttriads(counter.counters);
  netcounters::counter_isolates(counter.counters);
  netcounters::counter_ctriads(counter.counters);
  netcounters::counter_mutual(counter.counters);
  // Counting and printing the results
  std::vector< double > counts = counter.count_all();
  std::cout «
                         : " « counts[0] « std::endl «
    "Transitive triads : " « counts[U] « std::endl «
"Transitive triads : " « counts[1] « std::endl «
"Isolates : " « counts[2] « std::endl «
                         : " « counts[3] « std::endl « : " « counts[4] « std::endl;
    "C triads
    "Mutuals
  return 0;
Compiling this program using g++
g++ -std=c++11 -Wall -pedantic 08-counts.cpp -o counts && ./counts
Yields the following output:
Current view
        1
               1
   1,]
            1
   2,] .
                   . 1
   3,] . . . . . . 4,] 1 . 1 .
   5,]
  0,] .
1,] 1
           1
               1
  2,] 1 .
                      1
   3,] . . .
4,] 1 . 1
   5,] . . . .
Edges
Transitive triads : 3
Isolates
C triads
Mutuals
```

Features

Efficient memory usage

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

Documentation

More information can be found in the Doxygen website here and in the PDF version of the documentation here.

Code of Conduct

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

4 Main Page

Chapter 2

Module Index

2.1 Modules

Here is a list of all modules:

ounting	. 11
tatistical Models	. 11
EFMArray counters	. 12
hylo counters	. 22
hylo rules	29

6 Module Index

Chapter 3

Class Index

3.1 Class List

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

BArray< Cell_Type, Data_Type >
Baseline class for binary arrays
BArrayCell < Cell_Type, Data_Type >
BArrayCell_const< Cell_Type, Data_Type >
BArrayDense < Cell_Type, Data_Type >
Baseline class for binary arrays
BArrayDenseCell< Cell_Type, Data_Type >
BArrayDenseCell_const< Cell_Type, Data_Type >
BArrayDenseCol < Cell_Type, Data_Type >
BArrayDenseCol_const< Cell_Type, Data_Type >
BArrayDenseRow< Cell_Type, Data_Type >
BArrayDenseRow_const< Cell_Type, Data_Type >
BArrayRow< Cell_Type, Data_Type >
BArrayRow_const< Cell_Type, Data_Type >
BArrayVector< Cell_Type, Data_Type >
Row or column of a BArray 86
BArrayVector_const< Cell_Type, Data_Type >
Cell< Cell_Type >
Entries in BArray. For now, it only has two members:
Cell_const< Cell_Type >
ConstBArrayRowlter< Cell_Type, Data_Type >
Counter< Array_Type, Data_Type >
A counter function based on change statistics
Counters < Array_Type, Data_Type >
Vector of counters
DEFMCounterData
Data class used to store arbitrary uint or double vectors
DEFMData
Data class for DEFM arrays
Entries < Cell_Type >
A wrapper class to store source, target, val from a BArray object 118
Flock
A Flock is a group of Geese
FreqTable < T >
Frequency table of vectors

8 Class Index

Geese	
Annotated Phylo Model	128
Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >	
General framework for discrete exponential models. This class allows generating discrete expo-	
nential models in the form of a linear exponential model:	147
NetCounterData	
Data class used to store arbitrary uint or double vectors	162
NetworkData	
Data class for Networks	163
Node	
A single node for the model	165
NodeData	
Data definition for the PhyloArray class	171
PhyloCounterData	172
PhyloRuleDynData	175
PowerSet < Array_Type, Data_Rule_Type >	
Powerset of a binary array	177
Progress	
A simple progress bar	184
Rule < Array_Type, Data_Type >	
Rule for determining if a cell should be included in a sequence	185
Rules < Array_Type, Data_Type >	
Vector of objects of class Rule	187
StatsCounter< Array_Type, Data_Type >	
Count stats for a single Array	191
Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >	
Compute the support of sufficient statistics	194
vecHasher< T >	

Chapter 4

File Index

4.1 File List

Here is a list of all files with brief descriptions:

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

10 File Index

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

Chapter 5

Module Documentation

5.1 Counting

Classes

class DEFMData

Data class for DEFM arrays.

class NetworkData

Data class for Networks.

class NodeData

Data definition for the PhyloArray class.

class Counter< Array_Type, Data_Type >

A counter function based on change statistics.

5.1.1 Detailed Description

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

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

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

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

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

5.2 Statistical Models

Statistical models available in barry.

Classes

class Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >

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

class Geese

Annotated Phylo Model.

· class Flock

A Flock is a group of Geese.

5.2.1 Detailed Description

Statistical models available in barry.

5.3 **DEFMArray** counters

Counters for network models.

Functions

```
• template<typename Tarray = DEFMArray>
  void counter_ones (DEFMCounters< Tarray > *counters, int covar_index=-1)
     Prevalence of ones.
• template<typename Tarray = DEFMArray>
  void counter transition (DEFMCounters< Tarray > *counters, std::vector< size t > coords, int covar ←
  index=-1)
     Prevalence of ones.
• template<typename Tarray = DEFMArray>
  void counter_fixed_effect (DEFMCounters < Tarray > *counters, int covar_index, double k)
     Prevalence of ones.
template<typename Tnet = Network>
  void counter_edges (NetCounters< Tnet > *counters)
     Number of edges.
• template<typename Tnet = Network>
  void counter isolates (NetCounters< Tnet > *counters)
     Number of isolated vertices.

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

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

    template<> void counter_istar2 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter_ostar2 (NetCounters< Tnet > *counters)

    template<> void counter ostar2 (NetCounters< NetworkDense > *counters)

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

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

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

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

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

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

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

    template<> void counter odegree15 (NetCounters< NetworkDense > *counters)

    template<typename Tnet = Network>

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

    NETWORK_COUNTER (init_single_attr)

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

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

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

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

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

Rules for network models

Parameters

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

template < typename Tarray = DEFMArray > void rules_zerodiag (DEFMRules < Tarray > *rules)
 Number of edges.

5.3.1 Detailed Description

Counters for network models.

Parameters

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

5.3.2 Function Documentation

5.3.2.1 counter_absdiff()

Sum of absolute attribute difference between ego and alter.

Definition at line 910 of file network.hpp.

5.3.2.2 counter_ctriads() [1/2]

Definition at line 665 of file network.hpp.

5.3.2.3 counter_ctriads() [2/2]

Definition at line 610 of file network.hpp.

5.3.2.4 counter_degree()

Counts number of vertices with a given out-degree.

Definition at line 1328 of file network.hpp.

5.3.2.5 counter_density()

Definition at line 731 of file network.hpp.

5.3.2.6 counter_diff()

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

Definition at line 955 of file network.hpp.

5.3.2.7 counter_edges()

Number of edges.

Definition at line 152 of file network.hpp.

5.3.2.8 counter_fixed_effect()

Prevalence of ones.

Template Parameters

Tarray	
--------	--

Parameters

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

Definition at line 339 of file defm.hpp.

5.3.2.9 counter_idegree() [1/2]

Definition at line 1172 of file network.hpp.

5.3.2.10 counter_idegree() [2/2]

Counts number of vertices with a given in-degree.

Definition at line 1125 of file network.hpp.

5.3.2.11 counter_idegree15() [1/2]

Definition at line 787 of file network.hpp.

5.3.2.12 counter_idegree15() [2/2]

Definition at line 759 of file network.hpp.

5.3.2.13 counter_isolates() [1/2]

Definition at line 215 of file network.hpp.

5.3.2.14 counter_isolates() [2/2]

Number of isolated vertices.

Definition at line 175 of file network.hpp.

5.3.2.15 counter_istar2() [1/2]

Definition at line 338 of file network.hpp.

5.3.2.16 counter_istar2() [2/2]

Definition at line 312 of file network.hpp.

5.3.2.17 counter_mutual()

Number of mutual ties.

Definition at line 256 of file network.hpp.

5.3.2.18 counter_nodecov()

Definition at line 1068 of file network.hpp.

5.3.2.19 counter_nodeicov()

Definition at line 1018 of file network.hpp.

5.3.2.20 counter_nodematch()

Definition at line 1093 of file network.hpp.

5.3.2.21 counter_nodeocov()

Definition at line 1043 of file network.hpp.

5.3.2.22 counter_odegree() [1/2]

Definition at line 1273 of file network.hpp.

5.3.2.23 counter_odegree() [2/2]

Counts number of vertices with a given out-degree.

Definition at line 1225 of file network.hpp.

5.3.2.24 counter_odegree15() [1/2]

Definition at line 864 of file network.hpp.

5.3.2.25 counter_odegree15() [2/2]

Definition at line 836 of file network.hpp.

5.3.2.26 counter_ones()

Prevalence of ones.

Template Parameters

Tarray	
--------	--

Parameters

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

Definition at line 166 of file defm.hpp.

5.3.2.27 counter_ostar2() [1/2]

Definition at line 404 of file network.hpp.

5.3.2.28 counter_ostar2() [2/2]

Definition at line 376 of file network.hpp.

5.3.2.29 counter_transition()

Prevalence of ones.

Template Parameters

Tarray

Parameters

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

Definition at line 213 of file defm.hpp.

5.3.2.30 counter_ttriads() [1/2]

Definition at line 531 of file network.hpp.

5.3.2.31 counter_ttriads() [2/2]

Definition at line 441 of file network.hpp.

5.3.2.32 NETWORK_COUNTER()

Definition at line 999 of file network.hpp.

5.3.2.33 rules_zerodiag()

Number of edges.

Definition at line 374 of file defm.hpp.

5.4 Phylo counters

Counters for phylogenetic modeling.

Functions

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

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

k genes gain function nfun

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

Keeps track of how many pairs of genes preserve pseudostate.

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

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

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

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

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

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

 Co-evolution (joint gain or loss)
- void counter_longest (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)
 Longest branch mutates (either by gain or by loss)
- void counter_neofun (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

 Total number of neofunctionalization events.
- void counter_pairwise_neofun_singlefun (PhyloCounters *counters, uint nfunA, unsigned int duplication=DEFAULT_DUPLICATI
 Total number of neofunctionalization events sum_u sum_{w < u} [x(u,a)*(1 x(w,a)) + (1 x(u,a)) * x(w,a)] change stat: delta{x(u,a): 0->1} = 1 2 * x(w,a)
- void counter_neofun_a2b (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION Total number of neofunctionalization events.
- void counter_co_opt (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)
 Function co-opting.
- void counter_k_genes_changing (PhyloCounters *counters, unsigned int k, unsigned int duplication=DEFAULT_DUPLICATION)
 Indicator function. Equals to one if k genes changed and zero otherwise.
- void counter_less_than_p_prop_genes_changing (PhyloCounters *counters, double p, unsigned int duplication=DEFAULT DUPLICATION)
 - Indicator function. Equals to one if k genes changed and zero otherwise.
- void counter_gains_from_0 (PhyloCounters *counters, std::vector < uint > nfun, unsigned int duplication=DEFAULT_DUPLICAT
 Used when all the functions are in 0 (like the root node prob.)

5.4 Phylo counters 23

• void counter_overall_gains_from_0 (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

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

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

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

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

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

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

5.4.1 Detailed Description

Counters for phylogenetic modeling.

Parameters

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

5.4.2 Function Documentation

5.4.2.1 counter_co_opt()

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 1393 of file phylo.hpp.

5.4.2.2 counter_cogain()

Co-evolution (joint gain or loss)

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

Definition at line 888 of file phylo.hpp.

5.4.2.3 counter_gains()

Functional gains for a specific function (nfun).

Definition at line 193 of file phylo.hpp.

5.4.2.4 counter_gains_from_0()

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

Needs to specify function a.

Definition at line 1727 of file phylo.hpp.

5.4.2.5 counter_gains_k_offspring()

```
void counter_gains_k_offspring (  PhyloCounters * counters, \\ std::vector < uint > nfun, \\ uint k = 1u, \\ unsigned int duplication = DEFAULT_DUPLICATION ) [inline]
```

k genes gain function nfun

Definition at line 253 of file phylo.hpp.

5.4 Phylo counters 25

5.4.2.6 counter_genes_changing()

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

Definition at line 325 of file phylo.hpp.

5.4.2.7 counter_k_genes_changing()

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

Definition at line 1491 of file phylo.hpp.

5.4.2.8 counter_less_than_p_prop_genes_changing()

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

< How many genes diverge the parent

Definition at line 1611 of file phylo.hpp.

5.4.2.9 counter_longest()

Longest branch mutates (either by gain or by loss)

Definition at line 945 of file phylo.hpp.

5.4.2.10 counter_loss()

Total count of losses for an specific function.

Definition at line 688 of file phylo.hpp.

5.4.2.11 counter_maxfuns()

Cap the number of functions per gene.

Definition at line 626 of file phylo.hpp.

5.4.2.12 counter_neofun()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1115 of file phylo.hpp.

5.4.2.13 counter_neofun_a2b()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1260 of file phylo.hpp.

5.4 Phylo counters 27

5.4.2.14 counter_overall_changes()

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

Definition at line 740 of file phylo.hpp.

5.4.2.15 counter_overall_gains()

Overall functional gains.

Total number of gains (irrespective of the function).

Definition at line 155 of file phylo.hpp.

5.4.2.16 counter_overall_gains_from_0()

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

Needs to specify function a.

Definition at line 1793 of file phylo.hpp.

5.4.2.17 counter_overall_loss()

Overall functional loss.

Definition at line 578 of file phylo.hpp.

5.4.2.18 counter_pairwise_first_gain()

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

Needs to specify function a. sum $x(a)^3(1-x(b))^3 + x(b)^3(1-x(a))^3 + x(a)^3 + x(b)^3 + (1-x(a))^3 + (1-x(a))^3$ Definition at line 2045 of file phylo.hpp.

5.4.2.19 counter pairwise neofun singlefun()

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

Definition at line 1196 of file phylo.hpp.

5.4.2.20 counter_pairwise_overall_change()

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

Needs to specify function a.

Definition at line 1841 of file phylo.hpp.

5.4.2.21 counter_pairwise_preserving()

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

Needs to specify function a. sum $x(a)^3(1-x(b))^3 + x(b)^3(1-x(a))^3 + x(a)^3 + x(b)^3 + (1-x(a))^3 + (1-x(b))^3$ Definition at line 1906 of file phylo.hpp. 5.5 Phylo rules 29

5.4.2.22 counter_preserve_pseudogene()

Keeps track of how many pairs of genes preserve pseudostate.

Definition at line 394 of file phylo.hpp.

5.4.2.23 counter_prop_genes_changing()

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

Definition at line 476 of file phylo.hpp.

5.4.2.24 counter_subfun()

Total count of Sub-functionalization events.

It requires to specify data = {funA, funB}

Definition at line 799 of file phylo.hpp.

5.5 Phylo rules

Rules for phylogenetic modeling.

Classes

· class PhyloRuleDynData

Functions

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

5.5.1 Detailed Description

Rules for phylogenetic modeling.

Parameters

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

5.5.2 Function Documentation

5.5.2.1 rule_dyn_limit_changes()

Overall functional gains.

Parameters

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

Returns

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

Definition at line 2177 of file phylo.hpp.

Chapter 6

Namespace Documentation

6.1 barry Namespace Reference

barry: Your go-to motif accountant

Namespaces

counters

Tree class and Treelterator class.

6.1.1 Detailed Description

barry: Your go-to motif accountant

6.2 barry::counters Namespace Reference

Tree class and Treelterator class.

Namespaces

- defm
- network
- phylo

6.2.1 Detailed Description

Tree class and Treelterator class.

- 6.3 barry::counters::defm Namespace Reference
- 6.4 barry::counters::network Namespace Reference
- 6.5 barry::counters::phylo Namespace Reference

6.6 CHECK Namespace Reference

Integer constants used to specify which cell should be check.

Variables

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

6.6.1 Detailed Description

Integer constants used to specify which cell should be check.

6.6.2 Variable Documentation

6.6.2.1 BOTH

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

Definition at line 28 of file typedefs.hpp.

6.6.2.2 NONE

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

Definition at line 29 of file typedefs.hpp.

6.6.2.3 ONE

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

Definition at line 30 of file typedefs.hpp.

6.6.2.4 TWO

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

Definition at line 31 of file typedefs.hpp.

6.7 EXISTS Namespace Reference

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

Variables

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

6.7.1 Detailed Description

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

6.7.2 Variable Documentation

6.7.2.1 AS_ONE

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

Definition at line 46 of file typedefs.hpp.

6.7.2.2 AS_ZERO

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

Definition at line 45 of file typedefs.hpp.

6.7.2.3 BOTH

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

Definition at line 39 of file typedefs.hpp.

6.7.2.4 NONE

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

Definition at line 40 of file typedefs.hpp.

6.7.2.5 ONE

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

Definition at line 41 of file typedefs.hpp.

6.7.2.6 TWO

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

Definition at line 42 of file typedefs.hpp.

6.7.2.7 UKNOWN

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

Definition at line 44 of file typedefs.hpp.

Chapter 7

Class Documentation

7.1 BArray < Cell_Type, Data_Type > Class Template Reference

Baseline class for binary arrays.

#include <barray-bones.hpp>

Public Member Functions

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

Get the edgelist.

- void transpose ()
- void clear (bool hard=true)
- void resize (uint N_, uint M_)
- void reserve ()
- void print (const char *fmt=nullptr,...) const
- bool is_dense () const noexcept

Constructors

Parameters

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

Generated by Doxygen

• BArray ()

Zero-size array.

BArray (uint N_, uint M_)

Empty array.

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

Edgelist with data.

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

Edgelist with no data (simpler)

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

Move operator.

- BArray< Cell_Type, Data_Type > & operator= (BArray< Cell_Type, Data_Type > &&x) noexcept
 Move assignment.
- void set_data (Data_Type *data_, bool delete_data_=false)

Set the data object.

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

Cell-wise insertion/deletion

Parameters

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

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

Column/row wise interchange

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

Arithmetic operators

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

Public Attributes

· bool visited = false

Friends

- class BArrayCell
 Cell Type, Data Type
- class BArrayCell_const< Cell_Type, Data_Type >

7.1.1 Detailed Description

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

Baseline class for binary arrays.

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

Template Parameters

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

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

7.1.2 Constructor & Destructor Documentation

7.1.2.1 BArray() [1/6]

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

Zero-size array.

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

7.1.2.2 BArray() [2/6]

Empty array.

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

7.1.2.3 BArray() [3/6]

Edgelist with data.

7.1.2.4 BArray() [4/6]

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

Edgelist with no data (simpler)

7.1.2.5 BArray() [5/6]

Copy constructor.

7.1.2.6 BArray() [6/6]

Move operator.

7.1.2.7 \sim BArray()

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

7.1.3 Member Function Documentation

7.1.3.1 clear()

7.1.3.2 col()

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

7.1.3.3 D() [1/2]

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

7.1.3.4 D() [2/2]

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

7.1.3.5 **D_ptr()** [1/2]

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

7.1.3.6 **D_ptr()** [2/2]

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

7.1.3.7 default_val()

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

7.1.3.8 flush_data()

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

7.1.3.9 get_cell()

7.1.3.10 get_col_vec() [1/2]

7.1.3.11 get_col_vec() [2/2]

7.1.3.12 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 >
```

7.1.3.13 get_row_vec() [1/2]

7.1.3.14 get_row_vec() [2/2]

7.1.3.15 insert_cell() [1/3]

7.1.3.16 insert_cell() [2/3]

7.1.3.17 insert_cell() [3/3]

7.1.3.18 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 234 of file barray-bones.hpp.

7.1.3.19 is_empty()

7.1.3.20 ncol()

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

7.1.3.21 nnozero()

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

7.1.3.22 nrow()

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

7.1.3.23 operator()() [1/2]

7.1.3.24 operator()() [2/2]

7.1.3.25 operator*=()

7.1.3.26 operator+=() [1/3]

7.1.3.27 operator+=() [2/3]

7.1.3.28 operator+=() [3/3]

7.1.3.29 operator-=() [1/3]

7.1.3.30 operator-=() [2/3]

7.1.3.31 operator-=() [3/3]

7.1.3.32 operator/=()

7.1.3.33 operator=() [1/2]

Move assignment.

7.1.3.34 operator=() [2/2]

Assignment constructor.

7.1.3.35 operator==()

7.1.3.36 out_of_range()

7.1.3.37 print()

7.1.3.38 reserve()

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

7.1.3.39 resize()

7.1.3.40 rm_cell()

7.1.3.41 row()

7.1.3.42 set_data()

Set the data object.

Parameters

data_	
delete_←	
data_	

7.1.3.43 swap_cells()

7.1.3.44 swap_cols()

7.1.3.45 swap_rows()

7.1.3.46 toggle_cell()

7.1.3.47 toggle_lock()

7.1.3.48 transpose()

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

7.1.3.49 zero_col()

7.1.3.50 zero_row()

7.1.4 Friends And Related Function Documentation

7.1.4.1 BArrayCell< Cell_Type, Data_Type >

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

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

7.1.4.2 BArrayCell_const < Cell_Type, Data_Type >

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

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

7.1.5 Member Data Documentation

7.1.5.1 visited

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

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

Definition at line 48 of file barray-bones.hpp.

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

• include/barry/barray-bones.hpp

7.2 BArrayCell< Cell_Type, Data_Type > Class Template Reference

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

Public Member Functions

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

7.2.1 Detailed Description

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

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

7.2.2 Constructor & Destructor Documentation

7.2.2.1 BArrayCell()

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

7.2.2.2 ∼BArrayCell()

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

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

7.2.3 Member Function Documentation

7.2.3.1 operator Cell_Type()

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

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

7.2.3.2 operator*=()

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

7.2.3.3 operator+=()

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

7.2.3.4 operator-=()

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

7.2.3.5 operator/=()

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

7.2.3.6 operator=()

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

7.2.3.7 operator==()

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

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

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

7.3 BArrayCell_const< Cell_Type, Data_Type > Class Template Reference

#include <barraycell-bones.hpp>

Public Member Functions

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

7.3.1 Detailed Description

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

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

7.3.2 Constructor & Destructor Documentation

7.3.2.1 BArrayCell const()

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

7.3.2.2 ~BArrayCell_const()

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

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

7.3.3 Member Function Documentation

7.3.3.1 operator Cell_Type()

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

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

7.3.3.2 operator"!=()

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

7.3.3.3 operator<()

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

7.3.3.4 operator<=()

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

7.3.3.5 operator==()

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

7.3.3.6 operator>()

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

7.3.3.7 operator>=()

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

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

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

7.4 BArrayDense< Cell_Type, Data_Type > Class Template Reference

Baseline class for binary arrays.

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

Public Member Functions

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

Get the edgelist.

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

Constructors

Parameters

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

• BArrayDense ()

Zero-size array.

BArrayDense (uint N_, uint M_)

Empty array.

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

Edgelist with data.

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

Edgelist with no data (simpler)

BArrayDense (const BArrayDense < Cell_Type, Data_Type > &Array_, bool copy_data=false)

Copy constructor.

BArrayDense< Cell_Type, Data_Type > & operator= (const BArrayDense< Cell_Type, Data_Type > &Array_)

Assignment constructor.

BArrayDense (BArrayDense < Cell_Type, Data_Type > &&x) noexcept

Move operator.

BArrayDense< Cell_Type, Data_Type > & operator= (BArrayDense< Cell_Type, Data_Type > &&x)
 noexcept

Move assignment.

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

Set the data object.

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

Cell-wise insertion/deletion

Parameters

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

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

Column/row wise interchange

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

Arithmetic operators

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

Public Attributes

bool visited = false

Friends

- class BArrayDenseCell
 Cell Type, Data Type
- class BArrayDenseCol< Cell Type, Data Type >
- class BArrayDenseCol const< Cell Type, Data Type >
- class BArrayDenseRow
 Cell Type, Data Type
- class BArrayDenseRow const< Cell Type, Data Type >

7.4.1 Detailed Description

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

Baseline class for binary arrays.

BArrayDense class objects are arbitrary 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.

7.4.2 Constructor & Destructor Documentation

7.4.2.1 BArrayDense() [1/6]

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

Zero-size array.

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

7.4.2.2 BArrayDense() [2/6]

Empty array.

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

7.4.2.3 BArrayDense() [3/6]

Edgelist with data.

7.4.2.4 BArrayDense() [4/6]

Edgelist with no data (simpler)

7.4.2.5 BArrayDense() [5/6]

Copy constructor.

7.4.2.6 BArrayDense() [6/6]

Move operator.

7.4.2.7 ∼BArrayDense()

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

7.4.3 Member Function Documentation

7.4.3.1 clear()

7.4.3.2 col() [1/2]

Definition at line 490 of file barraydense-meat.hpp.

7.4.3.3 col() [2/2]

Definition at line 476 of file barraydense-meat.hpp.

7.4.3.4 colsum()

7.4.3.5 D() [1/2]

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

7.4.3.6 D() [2/2]

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

7.4.3.7 **D_ptr()** [1/2]

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

7.4.3.8 D_ptr() [2/2]

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

7.4.3.9 default_val()

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

7.4.3.10 get_cell()

7.4.3.11 get_col_vec() [1/2]

7.4.3.12 get_col_vec() [2/2]

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

7.4.3.13 get_data()

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

7.4.3.14 get_entries()

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

Get the edgelist.

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

Returns

Entries<Cell_Type>

7.4.3.15 get_row_vec() [1/2]

7.4.3.16 get_row_vec() [2/2]

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

7.4.3.17 insert_cell() [1/2]

7.4.3.18 insert_cell() [2/2]

7.4.3.19 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 255 of file barraydense-bones.hpp.

7.4.3.20 is_empty()

7.4.3.21 ncol()

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

7.4.3.22 nnozero()

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

7.4.3.23 nrow()

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

7.4.3.24 operator()() [1/2]

7.4.3.25 operator()() [2/2]

7.4.3.26 operator*=()

7.4.3.27 operator+=() [1/3]

7.4.3.28 operator+=() [2/3]

7.4.3.29 operator+=() [3/3]

7.4.3.30 operator-=() [1/3]

7.4.3.31 operator-=() [2/3]

7.4.3.32 operator-=() [3/3]

7.4.3.33 operator/=()

7.4.3.34 operator=() [1/2]

Move assignment.

7.4.3.35 operator=() [2/2]

Assignment constructor.

7.4.3.36 operator==()

7.4.3.37 out_of_range()

7.4.3.38 print()

7.4.3.39 reserve()

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

7.4.3.40 resize()

7.4.3.41 rm_cell()

7.4.3.42 row() [1/2]

7.4.3.43 row() [2/2]

7.4.3.44 rowsum()

7.4.3.45 set_data()

Set the data object.

Parameters

data_	
delete_ <i>←</i>	
data_	

7.4.3.46 swap_cells()

```
uint j0,
uint i1,
uint j1,
bool check_bounds = true,
int check_exists = CHECK::BOTH,
int * report = nullptr )
```

7.4.3.47 swap_cols()

7.4.3.48 swap_rows()

7.4.3.49 toggle_cell()

7.4.3.50 toggle_lock()

7.4.3.51 transpose()

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

7.4.3.52 zero_col()

7.4.3.53 zero_row()

7.4.4 Friends And Related Function Documentation

7.4.4.1 BArrayDenseCell< Cell_Type, Data_Type >

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

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

7.4.4.2 BArrayDenseCol < Cell_Type, Data_Type >

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

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

7.4.4.3 BArrayDenseCol_const< Cell_Type, Data_Type >

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

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

7.4.4.4 BArrayDenseRow< Cell_Type, Data_Type>

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

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

7.4.4.5 BArrayDenseRow_const< Cell_Type, Data_Type >

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

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

7.4.5 Member Data Documentation

7.4.5.1 visited

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

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

Definition at line 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

7.5 BArrayDenseCell< Cell_Type, Data_Type > Class Template Reference

#include <barraydensecell-bones.hpp>

Public Member Functions

```
\bullet \ \ \mathsf{BArrayDenseCell} \ (\mathsf{BArrayDense} < \mathsf{Cell\_Type}, \ \mathsf{Data\_Type} > *\mathsf{Array\_}, \ \mathsf{uint} \ \mathsf{i\_}, \ \mathsf{bool} \ \mathsf{check\_bounds} = \mathsf{true})
```

- \sim BArrayDenseCell ()
- void operator= (const Cell_Type &val)
- void operator+= (const Cell_Type &val)
- void operator-= (const Cell_Type &val)
- void operator*= (const Cell_Type &val)
- void operator/= (const Cell_Type &val)
- operator Cell_Type () const
- bool operator== (const Cell_Type &val) const

Friends

- class BArrayDense< Cell_Type, Data_Type >
- class BArrayDenseCol< Cell Type, Data Type >
- class BArrayDenseCol_const< Cell_Type, Data_Type >

7.5.1 Detailed Description

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

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

7.5.2 Constructor & Destructor Documentation

7.5.2.1 BArrayDenseCell()

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

7.5.2.2 ~BArrayDenseCell()

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

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

7.5.3 Member Function Documentation

7.5.3.1 operator Cell_Type()

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

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

7.5.3.2 operator*=()

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

7.5.3.3 operator+=()

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

7.5.3.4 operator-=()

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

7.5.3.5 operator/=()

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

7.5.3.6 operator=()

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

7.5.3.7 operator==()

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

7.5.4 Friends And Related Function Documentation

7.5.4.1 BArrayDense < Cell_Type, Data_Type >

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

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

7.5.4.2 BArrayDenseCol < Cell_Type, Data_Type >

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

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

7.5.4.3 BArrayDenseCol_const< Cell_Type, Data_Type >

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

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

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

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

7.6 BArrayDenseCell_const< Cell_Type, Data_Type > Class Template Reference

7.6.1 Detailed Description

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

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

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

• include/barry/barraydense-bones.hpp

7.7 BArrayDenseCol< Cell_Type, Data_Type > Class Template Reference

#include <barraydensecol-bones.hpp>

Public Member Functions

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

Friends

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

7.7.1 Detailed Description

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

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

7.7.2 Constructor & Destructor Documentation

7.7.2.1 BArrayDenseCol()

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

7.7.3 Member Function Documentation

7.7.3.1 begin()

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

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

7.7.3.2 end()

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

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

7.7.3.3 operator()()

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

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

7.7.3.4 size()

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

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

7.7.4 Friends And Related Function Documentation

7.7.4.1 BArrayDense < Cell_Type, Data_Type >

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

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

7.7.4.2 BArrayDenseCell< Cell_Type, Data_Type >

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

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

7.7.4.3 BArrayDenseCell_const< Cell_Type, Data_Type >

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

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

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

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

7.8 BArrayDenseCol_const< Cell_Type, Data_Type > Class Template Reference

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

Public Member Functions

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

Friends

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

7.8.1 Detailed Description

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

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

7.8.2 Constructor & Destructor Documentation

7.8.2.1 BArrayDenseCol_const()

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

7.8.3 Member Function Documentation

7.8.3.1 begin()

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

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

7.8.3.2 end()

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

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

7.8.3.3 operator()()

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

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

7.8.3.4 size()

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

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

7.8.4 Friends And Related Function Documentation

7.8.4.1 BArrayDenseCell< Cell_Type, Data_Type >

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

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

7.8.4.2 BArrayDenseCell_const< Cell_Type, Data_Type >

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

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

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

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

7.9 BArrayDenseRow< Cell_Type, Data_Type > Class Template Reference

#include <barraydenserow-bones.hpp>

Public Member Functions

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

Friends

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

7.9.1 Detailed Description

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

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

7.9.2 Constructor & Destructor Documentation

7.9.2.1 BArrayDenseRow()

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

7.9.3 Member Function Documentation

7.9.3.1 begin()

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

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

7.9.3.2 end()

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

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

7.9.3.3 operator()()

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

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

7.9.3.4 size()

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

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

7.9.4 Friends And Related Function Documentation

7.9.4.1 BArrayDense< Cell_Type, Data_Type>

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

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

7.9.4.2 BArrayDenseCell< Cell_Type, Data_Type >

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

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

7.9.4.3 BArrayDenseCell_const< Cell_Type, Data_Type >

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

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

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

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

7.10 BArrayDenseRow_const< Cell_Type, Data_Type > Class Template Reference

#include <barraydenserow-bones.hpp>

Public Member Functions

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

Friends

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

7.10.1 Detailed Description

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

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

7.10.2 Constructor & Destructor Documentation

7.10.2.1 BArrayDenseRow_const()

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

7.10.3 Member Function Documentation

7.10.3.1 begin()

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

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

7.10.3.2 end()

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

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

7.10.3.3 operator()()

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

7.10.3.4 size()

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

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

7.10.4 Friends And Related Function Documentation

7.10.4.1 BArrayDenseCell < Cell_Type, Data_Type >

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

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

7.10.4.2 BArrayDenseCell_const < Cell_Type, Data_Type >

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

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

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

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

7.11 BArrayRow < Cell_Type, Data_Type > Class Template Reference

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

Public Member Functions

- BArrayRow (BArray< Cell_Type, Data_Type > *Array_, uint i_, bool check_bounds=true)
- ∼BArrayRow ()
- void operator= (const BArrayRow< Cell Type, Data Type > &val)
- void operator+= (const BArrayRow< Cell_Type, Data_Type > &val)
- void operator-= (const BArrayRow< Cell_Type, Data_Type > &val)
- $\bullet \ \ \mathsf{void} \ \mathsf{operator} *= (\mathsf{const} \ \mathsf{BArrayRow} < \mathsf{Cell_Type}, \ \mathsf{Data_Type} > \mathsf{\&val})$
- void operator/= (const BArrayRow< Cell_Type, Data_Type > &val)
- operator BArrayRow< Cell_Type, Data_Type > () const
- bool operator== (const BArrayRow< Cell_Type, Data_Type > &val) const

7.11.1 Detailed Description

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

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

7.11.2 Constructor & Destructor Documentation

7.11.2.1 BArrayRow()

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

7.11.2.2 ∼BArrayRow()

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

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

7.11.3 Member Function Documentation

7.11.3.1 operator BArrayRow< Cell_Type, Data_Type >()

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

7.11.3.2 operator*=()

7.11.3.3 operator+=()

7.11.3.4 operator-=()

7.11.3.5 operator/=()

7.11.3.6 operator=()

7.11.3.7 operator==()

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

• include/barry/barrayrow-bones.hpp

7.12 BArrayRow_const< Cell_Type, Data_Type > Class Template Reference

#include <barrayrow-bones.hpp>

Public Member Functions

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

7.12.1 Detailed Description

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

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

7.12.2 Constructor & Destructor Documentation

7.12.2.1 BArrayRow_const()

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

7.12.2.2 ~BArrayRow_const()

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

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

7.12.3 Member Function Documentation

7.12.3.1 operator BArrayRow_const< Cell_Type, Data_Type >()

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

7.12.3.2 operator"!=()

7.12.3.3 operator<()

7.12.3.4 operator<=()

7.12.3.5 operator==()

7.12.3.6 operator>()

7.12.3.7 operator>=()

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

• include/barry/barrayrow-bones.hpp

7.13 BArrayVector< Cell Type, Data Type > Class Template Reference

Row or column of a BArray

#include <barrayvector-bones.hpp>

Public Member Functions

- BArrayVector (BArray < Cell_Type, Data_Type > *Array_, uint &dim_ uint &i_, bool check_bounds=true)

 Construct a new BArrayVector object.
- ∼BArrayVector ()
- bool is_row () const noexcept
- bool is_col () const noexcept
- uint size () const noexcept
- std::vector< Cell_Type >::const_iterator begin () noexcept
- std::vector< Cell_Type >::const_iterator end () noexcept
- void operator= (const Cell_Type &val)
- void operator+= (const Cell_Type &val)
- void operator-= (const Cell_Type &val)
- void operator*= (const Cell Type &val)
- void operator/= (const Cell_Type &val)
- operator std::vector< Cell_Type > () const
- bool operator== (const Cell_Type &val) const

7.13.1 Detailed Description

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

Row or column of a BArray

Template Parameters

Cell_Type	
Data_Type	

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

7.13.2 Constructor & Destructor Documentation

7.13.2.1 BArrayVector()

Construct a new BArrayVector object.

Parameters

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

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

7.13.2.2 ~BArrayVector()

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

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

7.13.3 Member Function Documentation

7.13.3.1 begin()

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

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

7.13.3.2 end()

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

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

7.13.3.3 is_col()

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

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

7.13.3.4 is_row()

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

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

7.13.3.5 operator std::vector< Cell_Type >()

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

Definition at line 177 of file barrayvector-meat.hpp.

7.13.3.6 operator*=()

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

7.13.3.7 operator+=()

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

7.13.3.8 operator-=()

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

7.13.3.9 operator/=()

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

7.13.3.10 operator=()

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

7.13.3.11 operator==()

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

7.13.3.12 size()

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

Definition at line 41 of file barrayvector-meat.hpp.

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

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

7.14 BArrayVector_const< Cell_Type, Data_Type > Class Template Reference

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

Public Member Functions

- BArrayVector_const (const BArray< Cell_Type, Data_Type > *Array_, uint &dim_ uint &i_, bool check_bounds=true)
- ∼BArrayVector_const ()
- bool is_row () const noexcept
- bool is_col () const noexcept
- uint size () const noexcept
- std::vector< Cell Type >::const iterator begin () noexcept
- std::vector< Cell_Type >::const_iterator end () noexcept
- operator std::vector< Cell Type > () const
- bool operator== (const Cell_Type &val) const
- bool operator!= (const Cell_Type &val) const
- bool operator< (const Cell_Type &val) const
- bool operator> (const Cell_Type &val) const
- bool operator<= (const Cell_Type &val) const
- bool operator>= (const Cell Type &val) const

7.14.1 Detailed Description

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

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

7.14.2 Constructor & Destructor Documentation

7.14.2.1 BArrayVector_const()

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

7.14.2.2 ~BArrayVector_const()

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

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

7.14.3 Member Function Documentation

7.14.3.1 begin()

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

7.14.3.2 end()

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

7.14.3.3 is_col()

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

7.14.3.4 is_row()

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

7.14.3.5 operator std::vector< Cell_Type >()

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

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

7.14.3.6 operator"!=()

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

7.14.3.7 operator<()

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

7.14.3.8 operator<=()

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

7.14.3.9 operator==()

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

7.14.3.10 operator>()

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

7.14.3.11 operator>=()

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

7.14.3.12 size()

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

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

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

7.15 Cell< Cell_Type > Class Template Reference

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

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

Public Member Functions

- Cell ()
- Cell (Cell_Type value_, bool visited_=false, bool active_=true)
- ∼Cell ()
- Cell (const Cell< Cell_Type > &arg)
- Cell< Cell_Type > & operator= (const Cell< Cell_Type > &other)
- Cell (Cell< Cell_Type > &&arg) noexcept
- Cell< Cell_Type > & operator= (Cell< Cell_Type > &&other) noexcept
- void add (Cell_Type x)
- operator Cell_Type () const
- bool operator== (const Cell< Cell_Type > &rhs) const
- bool operator!= (const Cell< Cell_Type > &rhs) const
- void add (double x)
- void add (unsigned int x)
- void add (int x)
- Cell ()
- Cell ()
- Cell ()

Public Attributes

- Cell_Type value
- bool visited
- · bool active

7.15.1 Detailed Description

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

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

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

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

7.15.2 Constructor & Destructor Documentation

7.15.2.1 Cell() [1/7]

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

7.15.2.2 Cell() [2/7]

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

7.15.2.3 ∼Cell()

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

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

7.15.2.4 Cell() [3/7]

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

7.15.2.5 Cell() [4/7]

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

7.15.2.6 Cell() [5/7]

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

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

7.15.2.7 Cell() [6/7]

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

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

7.15.2.8 Cell() [7/7]

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

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

7.15.3 Member Function Documentation

7.15.3.1 add() [1/4]

7.15.3.2 add() [2/4]

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

7.15.3.3 add() [3/4]

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

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

7.15.3.4 add() [4/4]

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

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

7.15.3.5 operator Cell_Type()

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

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

7.15.3.6 operator"!=()

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

7.15.3.7 operator=() [1/2]

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

7.15.3.8 operator=() [2/2]

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

7.15.3.9 operator==()

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

7.15.4 Member Data Documentation

7.15.4.1 active

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

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

7.15.4.2 value

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

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

7.15.4.3 visited

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

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

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

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

7.16 Cell_const < Cell_Type > Class Template Reference

7.16.1 Detailed Description

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

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

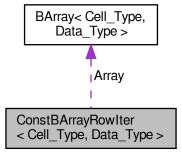
The documentation for this class was generated from the following file:

• include/barry/barray-meat.hpp

7.17 ConstBArrayRowlter< Cell_Type, Data_Type > Class Template Reference

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

Collaboration diagram for ConstBArrayRowlter< Cell_Type, Data_Type >:



Public Member Functions

- ConstBArrayRowlter (const BArray< Cell_Type, Data_Type > *Array_)
- ∼ConstBArrayRowIter ()

Public Attributes

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

7.17.1 Detailed Description

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

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

7.17.2 Constructor & Destructor Documentation

7.17.2.1 ConstBArrayRowIter()

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

7.17.2.2 ~ConstBArrayRowlter()

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

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

7.17.3 Member Data Documentation

7.17.3.1 Array

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

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

7.17.3.2 current_col

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

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

7.17.3.3 current_row

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

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

7.17.3.4 iter

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

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

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

• include/barry/barray-iterator.hpp

7.18 Counter< Array_Type, Data_Type > Class Template Reference

A counter function based on change statistics.

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

Public Member Functions

- ∼Counter ()
- double count (Array_Type &Array, uint i, uint j)
- double init (Array_Type &Array, uint i, uint j)
- std::string get_name () const
- std::string get_description () const

Creator passing a counter and an initializer

Parameters

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

- Counter ()
- Counter (Counter_fun_type< Array_Type, Data_Type > count_fun_, Counter_fun_type< Array_Type, Data_Type > init_fun_, Data_Type data_, std::string name_="", std::string desc_="")
- Counter (const Counter< Array_Type, Data_Type > &counter_)
 Copy constructor.
- Counter (Counter< Array_Type, Data_Type > &&counter_) noexcept Move constructor.
- Counter< Array_Type, Data_Type > operator= (const Counter< Array_Type, Data_Type > &counter_)
 Copy assignment.
- Counter< Array_Type, Data_Type > & operator= (Counter< Array_Type, Data_Type > &&counter_)
 noexcept

Move assignment.

Public Attributes

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

7.18.1 Detailed Description

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

A counter function based on change statistics.

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

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

7.18.2 Constructor & Destructor Documentation

7.18.2.1 Counter() [1/4]

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

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

7.18.2.2 Counter() [2/4]

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

7.18.2.3 Counter() [3/4]

Copy constructor.

7.18.2.4 Counter() [4/4]

Move constructor.

7.18.2.5 ∼Counter()

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

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

7.18.3 Member Function Documentation

7.18.3.1 count()

7.18.3.2 get_description()

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

7.18.3.3 get_name()

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

7.18.3.4 init()

7.18.3.5 operator=() [1/2]

Copy assignment.

7.18.3.6 operator=() [2/2]

Move assignment.

7.18.4 Member Data Documentation

7.18.4.1 count fun

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

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

7.18.4.2 data

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

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

7.18.4.3 desc

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

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

7.18.4.4 init_fun

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

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

7.18.4.5 name

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

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

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

• include/barry/counters-bones.hpp

7.19 Counters < Array_Type, Data_Type > Class Template Reference

Vector of counters.

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

Public Member Functions

- · Counters ()
- ∼Counters ()
- Counters (const Counters < Array_Type, Data_Type > &counter_)

Copy constructor.

Counters (Counters < Array_Type, Data_Type > &&counters_) noexcept

Move constructor.

- Counters< Array_Type, Data_Type > operator= (const Counters< Array_Type, Data_Type > &counter_)
 Copy assignment constructor.
- Counters< Array_Type, Data_Type > & operator= (Counters< Array_Type, Data_Type > &&counter_)
 noexcept

Move assignment constructor.

Counter< Array_Type, Data_Type > & operator[] (uint idx)

Returns a pointer to a particular counter.

• std::size_t size () const noexcept

Number of counters in the set.

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

7.19.1 Detailed Description

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

Vector of counters.

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

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

7.19.2 Constructor & Destructor Documentation

7.19.2.1 Counters() [1/3]

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

7.19.2.2 ∼Counters()

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

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

7.19.2.3 Counters() [2/3]

Copy constructor.

Parameters



7.19.2.4 Counters() [3/3]

Move constructor.

Parameters

counters⇔

7.19.3 Member Function Documentation

7.19.3.1 add_counter() [1/2]

7.19.3.2 add_counter() [2/2]

7.19.3.3 get_descriptions()

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

7.19.3.4 get_names()

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

7.19.3.5 operator=() [1/2]

Copy assignment constructor.

Parameters

counter←	

Returns

Counters<Array_Type,Data_Type>

7.19.3.6 operator=() [2/2]

Move assignment constructor.

Parameters



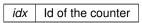
Returns

Counters<Array_Type,Data_Type>&

7.19.3.7 operator[]()

Returns a pointer to a particular counter.

Parameters



Returns

Counter<Array_Type,Data_Type>*

7.19.3.8 size()

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

Number of counters in the set.

Returns

uint

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

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

• include/barry/counters-bones.hpp

7.20 DEFMCounterData Class Reference

Data class used to store arbitrary uint or double vectors.

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

Public Member Functions

- DEFMCounterData ()
- DEFMCounterData (const std::vector< size_t > indices_, const std::vector< double > numbers_, size_t markov_order_)
- size_t idx (size_t i)
- double num (size_t i)
- ∼DEFMCounterData ()

Public Attributes

- std::vector< size t > indices
- std::vector< double > numbers
- · size_t markov_order

Order of the markov process.

7.20.1 Detailed Description

Data class used to store arbitrary uint or double vectors.

Definition at line 72 of file defm.hpp.

7.20.2 Constructor & Destructor Documentation

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

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

Definition at line 79 of file defm.hpp.

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

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

Definition at line 80 of file defm.hpp.

7.20.2.3 ~DEFMCounterData()

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

Definition at line 89 of file defm.hpp.

7.20.3 Member Function Documentation

7.20.3.1 idx()

Definition at line 86 of file defm.hpp.

7.20.3.2 num()

Definition at line 87 of file defm.hpp.

7.20.4 Member Data Documentation

7.20.4.1 indices

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

Definition at line 75 of file defm.hpp.

7.20.4.2 markov_order

```
size_t DEFMCounterData::markov_order
```

Order of the markov process.

Definition at line 77 of file defm.hpp.

7.20.4.3 numbers

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

Definition at line 76 of file defm.hpp.

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

• include/barry/counters/defm.hpp

7.21 DEFMData Class Reference

Data class for DEFM arrays.

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

Public Member Functions

- DEFMData ()
- DEFMData (std::vector< double > *covariates_, size_t obs_start_, size_t obs_n_times_, size_t n_← covariates_)

Constructor.

∼DEFMData ()

- 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

Public Attributes

std::vector< double > * covariates

Vector of covariates (complete vector)

size_t obs_start

Index of the observation in the data.

size_t obs_n_times

Number of records of the observation in the model.

size_t n_covariates

Number of covariates included in the model.

7.21.1 Detailed Description

Data class for DEFM arrays.

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

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

Definition at line 20 of file defm.hpp.

7.21.2 Constructor & Destructor Documentation

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

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

Definition at line 28 of file defm.hpp.

7.21.2.2 **DEFMData()** [2/2]

Constructor.

Parameters

covariates_	Pointer to the attribute data.
obs_start_	Location of the current observation in the covariates vector
obs_n_← times_	Number of observations in the model
Generated by Doxygen COVAriates_	Number of columns (covariates.)

Definition at line 38 of file defm.hpp.

7.21.2.3 ∼DEFMData()

```
\texttt{DEFMData::}{\sim} \texttt{DEFMData ( ) } \quad \texttt{[inline]}
```

Definition at line 56 of file defm.hpp.

7.21.3 Member Function Documentation

7.21.3.1 at()

Definition at line 65 of file defm.hpp.

7.21.3.2 operator()()

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

Parameters



Returns

double

Definition at line 60 of file defm.hpp.

7.21.4 Member Data Documentation

7.21.4.1 covariates

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

Vector of covariates (complete vector)

Definition at line 23 of file defm.hpp.

7.21.4.2 n_covariates

```
size_t DEFMData::n_covariates
```

Number of covariates included in the model.

Definition at line 26 of file defm.hpp.

7.21.4.3 obs_n_times

```
size_t DEFMData::obs_n_times
```

Number of records of the observation in the model.

Definition at line 25 of file defm.hpp.

7.21.4.4 obs_start

```
size_t DEFMData::obs_start
```

Index of the observation in the data.

Definition at line 24 of file defm.hpp.

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

• include/barry/counters/defm.hpp

7.22 Entries < Cell_Type > Class Template Reference

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

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

Public Member Functions

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

Public Attributes

```
• std::vector< uint > source
```

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

7.22.1 Detailed Description

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

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

Template Parameters

```
Cell_Type Any type
```

Definition at line 79 of file typedefs.hpp.

7.22.2 Constructor & Destructor Documentation

7.22.2.1 Entries() [1/2]

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

Definition at line 85 of file typedefs.hpp.

7.22.2.2 Entries() [2/2]

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

Definition at line 86 of file typedefs.hpp.

7.22.2.3 ∼Entries()

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

Definition at line 93 of file typedefs.hpp.

7.22.3 Member Function Documentation

7.22.3.1 resize()

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

Definition at line 95 of file typedefs.hpp.

7.22.4 Member Data Documentation

7.22.4.1 source

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

Definition at line 81 of file typedefs.hpp.

7.22.4.2 target

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

Definition at line 82 of file typedefs.hpp.

7.22.4.3 val

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

Definition at line 83 of file typedefs.hpp.

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

include/barry/typedefs.hpp

7.23 Flock Class Reference

A Flock is a group of Geese.

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

Public Member Functions

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

Add a tree to the flock.

· void set seed (const unsigned int &s)

Set the seed of the model.

- void init (unsigned int bar_width=BARRY_PROGRESS_BAR_WIDTH)
- phylocounters::PhyloCounters * get_counters ()
- phylocounters::PhyloSupport * get_support_fun ()
- std::vector< std::vector< double >> * get stats support ()
- std::vector< std::vector< double > > * get stats target ()
- phylocounters::PhyloModel * get_model ()

Returns the joint likelihood of the model.

• Geese * operator() (unsigned int i, bool check_bounds=true)

Access the i-th geese element.

Information about the model

- unsigned int nfuns () const noexcept
- unsigned int ntrees () const noexcept
- std::vector< unsigned int > nnodes () const noexcept
- std::vector< unsigned int > nleafs () const noexcept
- unsigned int nterms () const
- unsigned int support size () const noexcept
- std::vector< std::string > colnames () const
- unsigned int parse_polytomies (bool verb=true, std::vector< size_t > *dist=nullptr) const noexcept
 Check polytomies and return the largest.
- · void print () const

Public Attributes

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

7.23 Flock Class Reference 119

7.23.1 Detailed Description

A Flock is a group of Geese.

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

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

7.23.2 Constructor & Destructor Documentation

7.23.2.1 Flock()

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

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

7.23.2.2 ∼Flock()

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

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

7.23.3 Member Function Documentation

7.23.3.1 add data()

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

Add a tree to the flock.

Parameters

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

Returns

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

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

7.23.3.2 colnames()

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

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

7.23.3.3 get_counters()

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

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

7.23.3.4 get_model()

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

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

7.23.3.5 get_stats_support()

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

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

7.23.3.6 get_stats_target()

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

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

7.23 Flock Class Reference 121

7.23.3.7 get_support_fun()

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

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

7.23.3.8 init()

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

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

7.23.3.9 likelihood_joint()

Returns the joint likelihood of the model.

Parameters

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

7.23.3.10 nfuns()

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

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

7.23.3.11 nleafs()

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

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

7.23.3.12 nnodes()

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

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

7.23.3.13 nterms()

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

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

7.23.3.14 ntrees()

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

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

7.23.3.15 operator()()

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

Access the i-th geese element.

Parameters

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

Returns

Geese*

7.23 Flock Class Reference 123

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

7.23.3.16 parse_polytomies()

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

Check polytomies and return the largest.

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

7.23.3.17 print()

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

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

7.23.3.18 set seed()

Set the seed of the model.

Parameters

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

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

7.23.3.19 support_size()

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

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

7.23.4 Member Data Documentation

7.23.4.1 dat

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

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

7.23.4.2 initialized

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

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

7.23.4.3 model

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

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

7.23.4.4 nfunctions

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

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

7.23.4.5 rengine

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

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

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

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

7.24 FreqTable < T > Class Template Reference

Frequency table of vectors.

#include <freqtable.hpp>

Public Member Functions

```
FreqTable ()
~FreqTable ()
size_t add (const std::vector< T > &x, size_t *h_precomp)
Counts_type as_vector () const
const std::vector< double > & get_data () const
const std::unordered_map< size_t, size_t > & get_index () const
void clear ()
void reserve (size_t n, size_t k)
void print () const
size_t size () const noexcept
Number of unique elements in the table. (.
size_t make_hash (const std::vector< double > &x) const
```

7.24.1 Detailed Description

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

Frequency table of vectors.

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

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

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

Definition at line 27 of file freqtable.hpp.

7.24.2 Constructor & Destructor Documentation

7.24.2.1 FreqTable()

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

Definition at line 39 of file freqtable.hpp.

7.24.2.2 ∼FreqTable()

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

Definition at line 40 of file freqtable.hpp.

7.24.3 Member Function Documentation

7.24.3.1 add()

Definition at line 64 of file freqtable.hpp.

7.24.3.2 as_vector()

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

Definition at line 144 of file freqtable.hpp.

7.24.3.3 clear()

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

Definition at line 173 of file freqtable.hpp.

7.24.3.4 get_data()

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

Definition at line 45 of file freqtable.hpp.

7.24.3.5 get_index()

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

Definition at line 46 of file freqtable.hpp.

7.24.3.6 make_hash()

Definition at line 244 of file freqtable.hpp.

7.24.3.7 print()

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

Definition at line 209 of file freqtable.hpp.

7.24.3.8 reserve()

Definition at line 187 of file freqtable.hpp.

7.24.3.9 size()

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

Number of unique elements in the table. (.

Returns

size_t

Definition at line 236 of file freqtable.hpp.

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

include/barry/freqtable.hpp

7.25 Geese Class Reference

Annotated Phylo Model.

#include <defm-bones.hpp>

Public Member Functions

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

Prints information about the DEFM.

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

Powerset of a gene's possible states.

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

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

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

Prints information about the GEESE.

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

Powerset of a gene's possible states.

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

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

Construct a new Geese object

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

Parameters

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

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

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

Information about the model

Parameters

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

· unsigned int nfuns () const noexcept

Number of functions analyzed.

· unsigned int nnodes () const noexcept

Number of nodes (interior + leaf)

· unsigned int nleafs () const noexcept

Number of leaf.

• unsigned int nterms () const

Number of terms included.

• unsigned int support_size () const noexcept

Number of unique sets of sufficient stats.

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

Number of annotations.

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

Names of the terms in the model.

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

· unsigned int nfuns () const noexcept

Number of functions analyzed.

• unsigned int nnodes () const noexcept

Number of nodes (interior + leaf)

unsigned int nleafs () const noexcept

Number of leaf.

unsigned int nterms () const

Number of terms included.

unsigned int support_size () const noexcept

Number of unique sets of sufficient stats.

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

Number of annotations.

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

Names of the terms in the model.

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

Geese prediction

Calculate the conditional probability

Parameters

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

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

Returns

std::vector< double > Returns the posterior probability

- std::vector < std::vector < double > > predict_backend (const std::vector < double > &par, bool use_← reduced_sequence, const std::vector < uint > &preorder)
- std::vector< std::vector< double > > predict_exhaust_backend (const std::vector< double > &par, const std::vector< uint > &preorder)
- std::vector< std::vector< double > > predict_exhaust (const std::vector< double > &par)
- std::vector< std::vector< double > > predict_sim (const std::vector< double > &par, bool only_← annotated=false, unsigned int nsims=10000u)
- std::vector< std::vector< double >> predict (const std::vector< double > &par, std::vector< std::vector< double >> *res_prob=nullptr, bool leave_one_out=false, bool only_annotated=false, bool use_reduced
 _sequence=true)
- std::vector< std::vector< double > > predict_backend (const std::vector< double > &par, bool use_←
 reduced_sequence, const std::vector< uint > &preorder)
- std::vector< std::vector< double >> predict_exhaust_backend (const std::vector< double > &par, const std::vector< uint > &preorder)
- std::vector< std::vector< double > > predict_exhaust (const std::vector< double > &par)
- std::vector < std::vector < double > > predict_sim (const std::vector < double > &par, bool only_
 —
 annotated=false, unsigned int nsims=10000u)

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

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

Returns

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

std::mt19937 * get_rengine()
phylocounters::PhyloCounters * get_counters()
phylocounters::PhyloSupport * get_support_fun()
std::mt19937 * get_rengine()
phylocounters::PhyloCounters * get_counters()
phylocounters::PhyloCounters * get_support_fun()
phylocounters::PhyloModel * get_model()
phylocounters::PhyloSupport * get_support_fun()
```

Public Attributes

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

Locations of columns.

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

7.25.1 Detailed Description

Annotated Phylo Model.

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

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

7.25.2 Constructor & Destructor Documentation

7.25.2.1 Geese() [1/8]

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

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

7.25.2.2 Geese() [2/8]

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

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

7.25.2.3 Geese() [3/8]

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

7.25.2.4 Geese() [4/8]

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

7.25.2.5 ~Geese() [1/2]

```
Geese::∼Geese ( ) [inline]
```

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

7.25.2.6 Geese() [5/8]

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

133

7.25.2.7 Geese() [6/8]

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

7.25.2.8 Geese() [7/8]

7.25.2.9 Geese() [8/8]

7.25.2.10 ~Geese() [2/2]

```
Geese::∼Geese ( )
```

7.25.3 Member Function Documentation

7.25.3.1 calc_reduced_sequence() [1/2]

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

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

7.25.3.2 calc_reduced_sequence() [2/2]

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

7.25.3.3 calc_sequence() [1/2]

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

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

7.25.3.4 calc_sequence() [2/2]

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

7.25.3.5 colnames() [1/2]

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

Names of the terms in the model.

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

7.25.3.6 colnames() [2/2]

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

Names of the terms in the model.

7.25.3.7 get annotated nodes() [1/2]

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

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

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

7.25.3.8 get_annotated_nodes() [2/2]

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

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

7.25 Geese Class Reference 135

7.25.3.9 get_counters() [1/2]

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

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

7.25.3.10 get_counters() [2/2]

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

7.25.3.11 get_model() [1/2]

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

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

7.25.3.12 get_model() [2/2]

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

7.25.3.13 get_probabilities() [1/2]

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

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

7.25.3.14 get_probabilities() [2/2]

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

7.25.3.15 get_rengine() [1/2]

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

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

7.25.3.16 get_rengine() [2/2]

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

7.25.3.17 get_states() [1/2]

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

Powerset of a gene's possible states.

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

Returns

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

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

7.25.3.18 get_states() [2/2]

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

Powerset of a gene's possible states.

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

Returns

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

7.25.3.19 get_support_fun() [1/2]

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

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

7.25.3.20 get_support_fun() [2/2]

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

7.25 Geese Class Reference 137

7.25.3.21 inherit_support() [1/2]

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

7.25.3.22 inherit_support() [2/2]

7.25.3.23 init() [1/2]

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

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

7.25.3.24 init() [2/2]

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

7.25.3.25 init_node() [1/2]

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

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

7.25.3.26 init_node() [2/2]

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

7.25.3.27 likelihood() [1/2]

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

7.25.3.28 likelihood() [2/2]

7.25.3.29 likelihood exhaust() [1/2]

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

7.25.3.30 likelihood_exhaust() [2/2]

7.25.3.31 nannotations() [1/2]

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

Number of annotations.

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

7.25 Geese Class Reference 139

7.25.3.32 nannotations() [2/2]

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

Number of annotations.

7.25.3.33 nfuns() [1/2]

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

Number of functions analyzed.

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

7.25.3.34 nfuns() [2/2]

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

Number of functions analyzed.

7.25.3.35 nleafs() [1/2]

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

Number of leaf.

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

7.25.3.36 nleafs() [2/2]

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

Number of leaf.

7.25.3.37 nnodes() [1/2]

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

Number of nodes (interior + leaf)

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

7.25.3.38 nnodes() [2/2]

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

Number of nodes (interior + leaf)

7.25.3.39 nterms() [1/2]

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

Number of terms included.

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

7.25.3.40 nterms() [2/2]

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

Number of terms included.

7.25.3.41 observed_counts() [1/2]

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

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

7.25.3.42 observed_counts() [2/2]

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

7.25.3.43 operator=() [1/4]

7.25 Geese Class Reference 141

7.25.3.44 operator=() [2/4]

7.25.3.45 operator=() [3/4]

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

7.25.3.46 operator=() [4/4]

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

7.25.3.47 parse_polytomies() [1/2]

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

Check polytomies and return the largest.

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

7.25.3.48 parse_polytomies() [2/2]

Check polytomies and return the largest.

7.25.3.49 predict() [1/2]

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

7.25.3.50 predict() [2/2]

7.25.3.51 predict_backend() [1/2]

< True if the array belongs to the set

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

7.25.3.52 predict_backend() [2/2]

7.25.3.53 predict_exhaust() [1/2]

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

7.25.3.54 predict_exhaust() [2/2]

7.25 Geese Class Reference 143

7.25.3.55 predict_exhaust_backend() [1/2]

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

7.25.3.56 predict_exhaust_backend() [2/2]

7.25.3.57 predict_sim() [1/2]

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

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

7.25.3.58 predict_sim() [2/2]

7.25.3.59 print() [1/2]

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

Prints information about the DEFM.

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

7.25.3.60 print() [2/2]

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

Prints information about the GEESE.

7.25.3.61 print_observed_counts() [1/2]

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

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

7.25.3.62 print_observed_counts() [2/2]

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

7.25.3.63 set_seed() [1/2]

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

7.25.3.64 set_seed() [2/2]

7.25.3.65 simulate() [1/2]

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

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

7.25 Geese Class Reference 145

7.25.3.66 simulate() [2/2]

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

7.25.3.67 support_size() [1/2]

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

Number of unique sets of sufficient stats.

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

7.25.3.68 support_size() [2/2]

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

Number of unique sets of sufficient stats.

7.25.3.69 update_annotations() [1/2]

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

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

7.25.3.70 update_annotations() [2/2]

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

7.25.4 Member Data Documentation

7.25.4.1 delete_rengine

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

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

7.25.4.2 delete_support

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

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

7.25.4.3 initialized

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

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

7.25.4.4 map_to_nodes

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

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

7.25.4.5 nfunctions

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

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

7.25.4.6 nodes

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

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

7.25.4.7 pset_loc

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

Locations of columns.

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

7.25.4.8 reduced_sequence

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

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

7.25.4.9 sequence

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

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

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

- include/barry/models/defm/defm-bones.hpp
- include/barry/models/geese/geese-bones.hpp
- include/barry/models/defm/defm-meat.hpp
- include/barry/models/geese/geese-meat-constructors.hpp
- include/barry/models/geese/geese-meat-likelihood.hpp
- include/barry/models/geese/geese-meat-likelihood_exhaust.hpp
- include/barry/models/geese/geese-meat-predict.hpp
- include/barry/models/geese/geese-meat-predict_exhaust.hpp
- include/barry/models/geese/geese-meat-predict_sim.hpp
- include/barry/models/geese/geese-meat-simulate.hpp
- include/barry/models/geese/geese-meat.hpp

7.26 Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class Template Reference

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

#include <model-bones.hpp>

Public Member Functions

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

Adds an array to the support of not already included.

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

Prints information about the model.

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

Conditional probability ("Gibbs sampler")

- const std::mt19937 * get_rengine () const
- Counters < Array_Type, Data_Counter_Type > * get_counters ()
- Rules< Array_Type, Data_Rule_Type > * get_rules ()
- Rules< Array_Type, Data_Rule_Dyn_Type > * get_rules_dyn ()
- Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > * get_support_fun ()

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

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

- void add_counter (Counter< Array_Type, Data_Counter_Type > &counter)
- void add_counter (Counter_fun_type< Array_Type, Data_Counter_Type > count_fun_, Counter_fun_type
 Array Type, Data Counter Type > init fun =nullptr, Data Counter Type data =nullptr)
- void set_counters (Counters < Array_Type, Data_Counter_Type > *counters_)

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

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

- void add_rule (Rule < Array_Type, Data_Rule_Type > &rule)
- void add_rule (Rule_fun_type< Array_Type, Data_Rule_Type > count_fun_, Data_Rule_Type data_)
- void set_rules (Rules< Array_Type, Data_Rule_Type > *rules_)
- void add_rule_dyn (Rule< Array_Type, Data_Rule_Dyn_Type > &rule)
- void add_rule_dyn (Rule_fun_type< Array_Type, Data_Rule_Dyn_Type > count_fun_, Data_Rule_Dyn
 Type data)
- void set_rules_dyn (Rules < Array_Type, Data_Rule_Dyn_Type > *rules_)

Likelihood functions.

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

Parameters

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

- double likelihood (const std::vector< double > ¶ms, const uint &i, bool as_log=false)
- double likelihood (const std::vector < double > &target_, const uint &i, bool as log=false)
- double likelihood (const std::vector < double > ¶ms, const double *target_, const uint &i, bool as_←
 log=false)
- double likelihood total (const std::vector< double > ¶ms, bool as log=false)

Extract elements by index

Parameters

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

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

Size of the model

Number of different supports included in the model

This will return the size of stats_target.

Returns

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

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

Raw pointers to the support and target statistics.

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

Statistics of the support(s)

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

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

Set the transform_model_fun object.

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

7.26.1 Detailed Description

template<typename Array_Type = BArray<>>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>

class Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >

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

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

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

Template Parameters

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

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

7.26.2 Constructor & Destructor Documentation

7.26.2.1 Model() [1/3]

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

7.26.2.2 Model() [2/3]

7.26.2.3 Model() [3/3]

7.26.2.4 ~Model()

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

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

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

7.26.3 Member Function Documentation

7.26.3.1 add_array()

Adds an array to the support of not already included.

Parameters

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

Returns

The number of the array.

7.26.3.2 add_counter() [1/2]

7.26.3.3 add_counter() [2/2]

7.26.3.4 add_rule() [1/2]

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

7.26.3.5 add rule() [2/2]

7.26.3.6 add_rule_dyn() [1/2]

7.26.3.7 add_rule_dyn() [2/2]

7.26.3.8 colnames()

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

7.26.3.9 conditional_prob()

Conditional probability ("Gibbs sampler")

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

Parameters

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

Returns

double The conditional probability

7.26.3.10 gen_key()

7.26.3.11 get_arrays2support()

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

7.26.3.12 get_counters()

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

7.26.3.13 get_norm_const()

7.26.3.14 get pset()

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

7.26.3.15 get_pset_arrays()

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

7.26.3.16 get_pset_probs()

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

7.26.3.17 get_pset_stats() [1/2]

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

Statistics of the support(s)

7.26.3.18 get pset stats() [2/2]

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

7.26.3.19 get_rengine()

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

7.26.3.20 get_rules()

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

7.26.3.21 get_rules_dyn()

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

7.26.3.22 get_stats_support()

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

7.26.3.23 get_stats_target()

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

Raw pointers to the support and target statistics.

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

7.26.3.24 get support fun()

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

7.26.3.25 likelihood() [1/4]

7.26.3.26 likelihood() [2/4]

7.26.3.27 likelihood() [3/4]

7.26.3.28 likelihood() [4/4]

7.26.3.29 likelihood_total()

7.26.3.30 nterms()

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

7.26.3.31 operator=()

7.26.3.32 print()

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

Prints information about the model.

7.26.3.33 print_stats()

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

7.26.3.34 sample() [1/2]

7.26.3.35 sample() [2/2]

7.26.3.36 set_counters()

7.26.3.37 set_keygen()

7.26.3.38 set_rengine()

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

7.26.3.39 set_rules()

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

7.26.3.40 set_rules_dyn()

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

7.26.3.41 set_seed()

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

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

7.26.3.42 set_transform_model()

Set the transform_model_fun object.

The transform_model function is used to transform the data

Parameters

data	
target	
n_arrays	
arrays2support	

7.26.3.43 size()

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

7.26.3.44 size_unique()

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

7.26.3.45 store psets()

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

7.26.3.46 support_size()

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

7.26.3.47 transform_model()

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

• include/barry/model-bones.hpp

7.27 NetCounterData Class Reference

Data class used to store arbitrary uint or double vectors.

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

Public Member Functions

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

Public Attributes

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

7.27.1 Detailed Description

Data class used to store arbitrary uint or double vectors.

Definition at line 56 of file network.hpp.

7.27.2 Constructor & Destructor Documentation

7.27.2.1 NetCounterData() [1/2]

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

Definition at line 62 of file network.hpp.

7.27.2.2 NetCounterData() [2/2]

Definition at line 63 of file network.hpp.

7.27.2.3 ∼NetCounterData()

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

Definition at line 68 of file network.hpp.

7.27.3 Member Data Documentation

7.27.3.1 indices

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

Definition at line 59 of file network.hpp.

7.27.3.2 numbers

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

Definition at line 60 of file network.hpp.

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

• include/barry/counters/network.hpp

7.28 NetworkData Class Reference

Data class for Networks.

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

Public Member Functions

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

Constructor using a single attribute.

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

Constructor using multiple attributes.

∼NetworkData ()

Public Attributes

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

7.28.1 Detailed Description

Data class for Networks.

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

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

Definition at line 19 of file network.hpp.

7.28.2 Constructor & Destructor Documentation

7.28.2.1 NetworkData() [1/3]

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

Definition at line 25 of file network.hpp.

7.28.2.2 NetworkData() [2/3]

Constructor using a single attribute.

Parameters

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

Definition at line 33 of file network.hpp.

7.28.2.3 NetworkData() [3/3]

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

Constructor using multiple attributes.

7.29 Node Class Reference 165

Parameters

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

Definition at line 45 of file network.hpp.

7.28.2.4 ~NetworkData()

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

Definition at line 51 of file network.hpp.

7.28.3 Member Data Documentation

7.28.3.1 directed

bool NetworkData::directed = true

Definition at line 22 of file network.hpp.

7.28.3.2 vertex_attr

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

Definition at line 23 of file network.hpp.

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

• include/barry/counters/network.hpp

7.29 Node Class Reference

A single node for the model.

#include <geese-node-bones.hpp>

Collaboration diagram for Node:



Public Member Functions

- ∼Node ()
- int get_parent () const
- · unsigned int noffspring () const noexcept
- bool is_leaf () const noexcept

Construct a new Node object

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

Public Attributes

· unsigned int id

Id of the node (as specified in the input)

· unsigned int ord

Order in which the node was created.

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

Observed annotations (only defined for Geese)

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

Arrays given all possible states.

Node * parent = nullptr

Parent node.

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

Offspring nodes.

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

ID of the array in the model.

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

Induced subtree probabilities.

std::vector< double > probability

The probability of observing each state.

7.29.1 Detailed Description

A single node for the model.

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

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

7.29.2 Constructor & Destructor Documentation

7.29 Node Class Reference 167

7.29.2.1 Node() [1/5]

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

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

7.29.2.2 Node() [2/5]

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

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

7.29.2.3 Node() [3/5]

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

7.29.2.4 Node() [4/5]

```
Node::Node (
          Node && x ) [inline], [noexcept]
```

Definition at line 69 of file geese-node-bones.hpp.

7.29.2.5 Node() [5/5]

```
Node::Node ( {\tt const\ Node\ \&\ x\ )} \quad [{\tt inline}]
```

Definition at line 83 of file geese-node-bones.hpp.

7.29.2.6 ∼Node()

```
Node::~Node ( ) [inline]
```

Definition at line 47 of file geese-node-bones.hpp.

7.29.3 Member Function Documentation

7.29.3.1 get_parent()

```
int Node::get_parent ( ) const [inline]
```

Definition at line 97 of file geese-node-bones.hpp.

7.29.3.2 is_leaf()

```
bool Node::is_leaf ( ) const [inline], [noexcept]
```

Definition at line 109 of file geese-node-bones.hpp.

7.29.3.3 noffspring()

```
unsigned int Node::noffspring ( ) const [inline], [noexcept]
```

Definition at line 103 of file geese-node-bones.hpp.

7.29.4 Member Data Documentation

7.29.4.1 annotations

```
std::vector< unsigned int > Node::annotations
```

Observed annotations (only defined for Geese)

Definition at line 18 of file geese-node-bones.hpp.

7.29 Node Class Reference 169

7.29.4.2 array

phylocounters::PhyloArray Node::array

Definition at line 17 of file geese-node-bones.hpp.

7.29.4.3 arrays

```
std::vector< phylocounters::PhyloArray > Node::arrays = {}
```

Arrays given all possible states.

Definition at line 21 of file geese-node-bones.hpp.

7.29.4.4 duplication

bool Node::duplication

Definition at line 19 of file geese-node-bones.hpp.

7.29.4.5 id

unsigned int Node::id

Id of the node (as specified in the input)

Definition at line 14 of file geese-node-bones.hpp.

7.29.4.6 narray

```
std::vector< unsigned int > Node::narray = {}
```

ID of the array in the model.

Definition at line 24 of file geese-node-bones.hpp.

7.29.4.7 offspring

```
std::vector< Node* > Node::offspring = {}
```

Offspring nodes.

Definition at line 23 of file geese-node-bones.hpp.

7.29.4.8 ord

```
unsigned int Node::ord
```

Order in which the node was created.

Definition at line 15 of file geese-node-bones.hpp.

7.29.4.9 parent

```
Node* Node::parent = nullptr
```

Parent node.

Definition at line 22 of file geese-node-bones.hpp.

7.29.4.10 probability

```
std::vector< double > Node::probability
```

The probability of observing each state.

Definition at line 28 of file geese-node-bones.hpp.

7.29.4.11 subtree_prob

```
std::vector< double > Node::subtree_prob
```

Induced subtree probabilities.

Definition at line 27 of file geese-node-bones.hpp.

7.29.4.12 visited

```
bool Node::visited = false
```

Definition at line 25 of file geese-node-bones.hpp.

The documentation for this class was generated from the following file:

• include/barry/models/geese/geese-node-bones.hpp

7.30 NodeData Class Reference

Data definition for the PhyloArray class.

```
#include <phylo.hpp>
```

Public Member Functions

NodeData (const std::vector< double > &blengths_, const std::vector< bool > &states_, bool duplication
 —=true)

Public Attributes

```
std::vector< double > blengths = {}std::vector< bool > states = {}
```

• bool duplication = true

7.30.1 Detailed Description

Data definition for the PhyloArray class.

Details about the available counters for PhyloArray objects can be found in the Phylo counters section.

This holds basic information about a given node.

Definition at line 38 of file phylo.hpp.

7.30.2 Constructor & Destructor Documentation

7.30.2.1 NodeData()

Definition at line 58 of file phylo.hpp.

7.30.3 Member Data Documentation

7.30.3.1 blengths

```
std::vector< double > NodeData::blengths = {}
```

Branch length.

Definition at line 44 of file phylo.hpp.

7.30.3.2 duplication

```
bool NodeData::duplication = true
```

Definition at line 54 of file phylo.hpp.

7.30.3.3 states

```
std::vector< bool > NodeData::states = {}
```

State of the parent node.

Definition at line 49 of file phylo.hpp.

The documentation for this class was generated from the following file:

• include/barry/counters/phylo.hpp

7.31 PhyloCounterData Class Reference

```
#include <phylo.hpp>
```

Public Member Functions

- PhyloCounterData (std::vector< uint > data_, std::vector< double > *counters_=nullptr)
- PhyloCounterData ()
- uint at (uint d)
- uint operator() (uint d)
- uint operator[] (uint d)
- void reserve (uint x)
- void push_back (uint x)
- void shrink_to_fit ()
- uint size ()
- std::vector< uint >::iterator begin ()
- std::vector< uint >::iterator end ()
- bool empty ()
- std::vector< double > * get_counters ()

7.31.1 Detailed Description

Definition at line 69 of file phylo.hpp.

7.31.2 Constructor & Destructor Documentation

7.31.2.1 PhyloCounterData() [1/2]

Definition at line 75 of file phylo.hpp.

7.31.2.2 PhyloCounterData() [2/2]

```
PhyloCounterData::PhyloCounterData ( ) [inline]
```

Definition at line 80 of file phylo.hpp.

7.31.3 Member Function Documentation

7.31.3.1 at()

Definition at line 82 of file phylo.hpp.

7.31.3.2 begin()

```
std::vector< uint >::iterator PhyloCounterData::begin ( ) [inline]
```

Definition at line 90 of file phylo.hpp.

7.31.3.3 empty()

```
bool PhyloCounterData::empty ( ) [inline]
```

Definition at line 93 of file phylo.hpp.

7.31.3.4 end()

```
std::vector< uint >::iterator PhyloCounterData::end ( ) [inline]
```

Definition at line 91 of file phylo.hpp.

7.31.3.5 get_counters()

```
std::vector< double >* PhyloCounterData::get_counters ( ) [inline]
```

Definition at line 94 of file phylo.hpp.

7.31.3.6 operator()()

Definition at line 83 of file phylo.hpp.

7.31.3.7 operator[]()

```
uint PhyloCounterData::operator[] (
          uint d) [inline]
```

Definition at line 84 of file phylo.hpp.

7.31.3.8 push_back()

Definition at line 86 of file phylo.hpp.

7.31.3.9 reserve()

Definition at line 85 of file phylo.hpp.

7.31.3.10 shrink_to_fit()

```
void PhyloCounterData::shrink_to_fit ( ) [inline]
```

Definition at line 87 of file phylo.hpp.

7.31.3.11 size()

```
uint PhyloCounterData::size ( ) [inline]
```

Definition at line 88 of file phylo.hpp.

The documentation for this class was generated from the following file:

include/barry/counters/phylo.hpp

7.32 PhyloRuleDynData Class Reference

```
#include <phylo.hpp>
```

Public Member Functions

- PhyloRuleDynData (const std::vector< double > *counts_, uint pos_, uint lb_, uint ub_, uint duplication_)
- \sim PhyloRuleDynData ()

Public Attributes

- const std::vector< double > * counts
- uint pos
- uint lb
- uint ub
- · uint duplication

7.32.1 Detailed Description

Definition at line 2147 of file phylo.hpp.

7.32.2 Constructor & Destructor Documentation

7.32.2.1 PhyloRuleDynData()

Definition at line 2155 of file phylo.hpp.

7.32.2.2 ~PhyloRuleDynData()

```
PhyloRuleDynData::~PhyloRuleDynData ( ) [inline]
```

Definition at line 2164 of file phylo.hpp.

7.32.3 Member Data Documentation

7.32.3.1 counts

```
const std::vector< double >* PhyloRuleDynData::counts
```

Definition at line 2149 of file phylo.hpp.

7.32.3.2 duplication

```
uint PhyloRuleDynData::duplication
```

Definition at line 2153 of file phylo.hpp.

7.32.3.3 lb

```
uint PhyloRuleDynData::lb
```

Definition at line 2151 of file phylo.hpp.

7.32.3.4 pos

uint PhyloRuleDynData::pos

Definition at line 2150 of file phylo.hpp.

7.32.3.5 ub

uint PhyloRuleDynData::ub

Definition at line 2152 of file phylo.hpp.

The documentation for this class was generated from the following file:

• include/barry/counters/phylo.hpp

7.33 PowerSet< Array_Type, Data_Rule_Type > Class Template Reference

Powerset of a binary array.

#include <powerset-bones.hpp>

 $Collaboration\ diagram\ for\ PowerSet < Array_Type,\ Data_Rule_Type >:$



Public Member Functions

- void init support ()
- void calc ()
- void reset (uint N_, uint M_)

Construct and destroy a PowerSet object

- PowerSet ()
- PowerSet (uint N_, uint M_)
- PowerSet (const Array_Type & array)
- ∼PowerSet ()

Wrappers for the <tt>Rules</tt> member.

These will add rules to the model, which are shared by the support and the actual counter function.

- $\bullet \ \ \mathsf{void} \ \ \mathsf{add_rule} \ \ (\mathsf{Rule} {<} \ \mathsf{Array_Type}, \ \mathsf{Data_Rule_Type} > \mathsf{rule})$
- void add_rule (Rule_fun_type< Array_Type, Data_Rule_Type > count_fun_, Data_Rule_Type data_)

Getter functions

- const std::vector< Array_Type > * get_data_ptr () const
- std::vector< Array_Type > get_data () const
- std::vector< Array_Type >::iterator begin ()
- std::vector< Array_Type >::iterator end ()
- std::size_t size () const noexcept
- const Array_Type & operator[] (const unsigned int &i) const

Public Attributes

- Array_Type EmptyArray
- $std::vector < Array_Type > data$
- Rules
 Array_Type, Data_Rule_Type > * rules
- uint N
- uint M
- bool rules deleted = false
- std::vector < size_t > coordinates_free
- std::vector< size_t > coordinates_locked
- size_t n_free
- size_t n_locked

7.33.1 Detailed Description

template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool> class PowerSet< Array_Type, Data_Rule_Type >

Powerset of a binary array.

Template Parameters

Array_Type	
Data_Rule_Type	

Definition at line 17 of file powerset-bones.hpp.

7.33.2 Constructor & Destructor Documentation

7.33.2.1 PowerSet() [1/3]

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
PowerSet< Array_Type, Data_Rule_Type >::PowerSet ( ) [inline]
```

Definition at line 42 of file powerset-bones.hpp.

7.33.2.2 PowerSet() [2/3]

Definition at line 44 of file powerset-bones.hpp.

7.33.2.3 PowerSet() [3/3]

Definition at line 7 of file powerset-meat.hpp.

7.33.2.4 ∼PowerSet()

```
template<typename Array_Type , typename Data_Rule_Type >
PowerSet< Array_Type, Data_Rule_Type >::~PowerSet [inline]
```

Definition at line 15 of file powerset-meat.hpp.

7.33.3 Member Function Documentation

7.33.3.1 add_rule() [1/2]

Definition at line 175 of file powerset-meat.hpp.

7.33.3.2 add_rule() [2/2]

Definition at line 184 of file powerset-meat.hpp.

7.33.3.3 begin()

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< Array_Type >::iterator PowerSet< Array_Type, Data_Rule_Type >::begin ( ) [inline]
```

Definition at line 74 of file powerset-bones.hpp.

7.33.3.4 calc()

```
template<typename Array_Type , typename Data_Rule_Type >
void PowerSet< Array_Type, Data_Rule_Type >::calc [inline]
```

Definition at line 146 of file powerset-meat.hpp.

7.33.3.5 end()

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< Array_Type >::iterator PowerSet< Array_Type, Data_Rule_Type >::end ( ) [inline]
```

Definition at line 75 of file powerset-bones.hpp.

7.33.3.6 get_data()

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< Array_Type > PowerSet< Array_Type, Data_Rule_Type >::get_data ( ) const [inline]
```

Definition at line 73 of file powerset-bones.hpp.

7.33.3.7 get_data_ptr()

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
const std::vector< Array_Type >* PowerSet< Array_Type, Data_Rule_Type >::get_data_ptr ()
const [inline]
```

Definition at line 72 of file powerset-bones.hpp.

7.33.3.8 init support()

```
template<typename Array_Type , typename Data_Rule_Type >
void PowerSet< Array_Type, Data_Rule_Type >::init_support [inline]
```

Definition at line 21 of file powerset-meat.hpp.

7.33.3.9 operator[]()

Definition at line 77 of file powerset-bones.hpp.

7.33.3.10 reset()

Definition at line 162 of file powerset-meat.hpp.

7.33.3.11 size()

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::size_t PowerSet< Array_Type, Data_Rule_Type >::size ( ) const [inline], [noexcept]
```

Definition at line 76 of file powerset-bones.hpp.

7.33.4 Member Data Documentation

7.33.4.1 coordinates_free

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< size_t > PowerSet< Array_Type, Data_Rule_Type >::coordinates_free
```

Definition at line 32 of file powerset-bones.hpp.

7.33.4.2 coordinates locked

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< size_t > PowerSet< Array_Type, Data_Rule_Type >::coordinates_locked
```

Definition at line 33 of file powerset-bones.hpp.

7.33.4.3 data

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< Array_Type > PowerSet< Array_Type, Data_Rule_Type >::data
```

Definition at line 25 of file powerset-bones.hpp.

7.33.4.4 EmptyArray

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
Array_Type PowerSet< Array_Type, Data_Rule_Type >::EmptyArray
```

Definition at line 24 of file powerset-bones.hpp.

7.33.4.5 M

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
uint PowerSet< Array_Type, Data_Rule_Type >::M
```

Definition at line 28 of file powerset-bones.hpp.

7.33.4.6 N

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
uint PowerSet< Array_Type, Data_Rule_Type >::N
```

Definition at line 28 of file powerset-bones.hpp.

7.33.4.7 n_free

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
size_t PowerSet< Array_Type, Data_Rule_Type >::n_free
```

Definition at line 34 of file powerset-bones.hpp.

7.33.4.8 n locked

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
size_t PowerSet< Array_Type, Data_Rule_Type >::n_locked
```

Definition at line 35 of file powerset-bones.hpp.

7.33.4.9 rules

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
Rules<Array_Type,Data_Rule_Type>* PowerSet< Array_Type, Data_Rule_Type >::rules
```

Definition at line 26 of file powerset-bones.hpp.

7.33.4.10 rules_deleted

```
template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
bool PowerSet< Array_Type, Data_Rule_Type >::rules_deleted = false
```

Definition at line 29 of file powerset-bones.hpp.

The documentation for this class was generated from the following files:

- include/barry/powerset-bones.hpp
- include/barry/powerset-meat.hpp

7.34 Progress Class Reference

A simple progress bar.

```
#include progress.hpp>
```

Public Member Functions

```
• Progress (int n_, int width_)
```

- ∼Progress ()
- void next ()
- void end ()

7.34.1 Detailed Description

A simple progress bar.

Definition at line 11 of file progress.hpp.

7.34.2 Constructor & Destructor Documentation

7.34.2.1 Progress()

```
Progress::Progress (
          int n_,
          int width_ ) [inline]
```

Definition at line 30 of file progress.hpp.

7.34.2.2 ∼Progress()

```
Progress::~Progress ( ) [inline]
```

Definition at line 23 of file progress.hpp.

7.34.3 Member Function Documentation

7.34.3.1 end()

```
void Progress::end ( ) [inline]
```

Definition at line 52 of file progress.hpp.

7.34.3.2 next()

```
void Progress::next ( ) [inline]
```

Definition at line 41 of file progress.hpp.

The documentation for this class was generated from the following file:

• include/barry/progress.hpp

7.35 Rule < Array_Type, Data_Type > Class Template Reference

Rule for determining if a cell should be included in a sequence.

```
#include <rules-bones.hpp>
```

Public Member Functions

- \sim Rule ()
- Data_Type & D ()

Read/Write access to the data.

bool operator() (const Array_Type &a, uint i, uint j)

Construct a new Rule object

Construct a new Rule object

Parameters

fun_	A function of type Rule_fun_type.
dat_	Data pointer to be passed to fun_
delete_← dat_	When true, the Rule destructor will delete the pointer, if defined.

- Rule ()
- Rule (Rule_fun_type< Array_Type, Data_Type > fun_, Data_Type dat_)

7.35.1 Detailed Description

```
\label{template} $$ \ensuremath{\sf template}$$ < typename \ensuremath{\sf Array\_Type}$ = BArray<>, typename \ensuremath{\sf Data\_Type}$ = bool> class \ensuremath{\sf Rule}< Array\_Type, \ensuremath{\sf Data\_Type}>
```

Rule for determining if a cell should be included in a sequence.

Rules can be used together with Support and PowerSet to determine which cells should be included when enumerating all possible realizations of a binary array.

Template Parameters

Array_Type	An object of class BArray.
Data_Type	Any type.

Definition at line 22 of file rules-bones.hpp.

7.35.2 Constructor & Destructor Documentation

7.35.2.1 Rule() [1/2]

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Rule< Array_Type, Data_Type >::Rule ( ) [inline]
```

Definition at line 40 of file rules-bones.hpp.

7.35.2.2 Rule() [2/2]

Definition at line 41 of file rules-bones.hpp.

7.35.2.3 ∼Rule()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Rule< Array_Type, Data_Type >::~Rule ( ) [inline]
```

Definition at line 47 of file rules-bones.hpp.

7.35.3 Member Function Documentation

7.35.3.1 D()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Data_Type& Rule< Array_Type, Data_Type >::D ()
```

Read/Write access to the data.

7.35.3.2 operator()()

Definition at line 42 of file rules-meat.hpp.

The documentation for this class was generated from the following files:

- include/barry/rules-bones.hpp
- include/barry/rules-meat.hpp

7.36 Rules < Array_Type, Data_Type > Class Template Reference

Vector of objects of class Rule.

```
#include <rules-bones.hpp>
```

Public Member Functions

- Rules ()
- Rules (const Rules < Array_Type, Data_Type > &rules_)
- Rules
 Array_Type, Data_Type > operator= (const Rules
 Array_Type, Data_Type > &rules_)
- ∼Rules ()
- uint size () const noexcept
- bool operator() (const Array_Type &a, uint i, uint j)

Check whether a given cell is free or locked.

void get_seq (const Array_Type &a, std::vector< size_t > *free, std::vector< size_t > *locked=nullptr)
 Computes the sequence of free and locked cells in an BArray.

Rule adding

Parameters

```
rule
```

- void add_rule (Rule < Array_Type, Data_Type > rule)
- void add_rule (Rule_fun_type< Array_Type, Data_Type > rule_, Data_Type data_)

7.36.1 Detailed Description

```
template<typename Array_Type, typename Data_Type> class Rules< Array_Type, Data_Type >
```

Vector of objects of class Rule.

Template Parameters

Array_Type	An object of class BArray
Data_Type	Any type.

Definition at line 62 of file rules-bones.hpp.

7.36.2 Constructor & Destructor Documentation

7.36.2.1 Rules() [1/2]

```
template<typename Array_Type , typename Data_Type >
Rules< Array_Type, Data_Type >::Rules ( ) [inline]
```

Definition at line 68 of file rules-bones.hpp.

7.36.2.2 Rules() [2/2]

Definition at line 10 of file rules-meat.hpp.

7.36.2.3 ∼Rules()

```
template<typename Array_Type , typename Data_Type >
Rules< Array_Type, Data_Type >::~Rules ( ) [inline]
```

Definition at line 73 of file rules-bones.hpp.

7.36.3 Member Function Documentation

7.36.3.1 add_rule() [1/2]

Definition at line 47 of file rules-meat.hpp.

7.36.3.2 add_rule() [2/2]

Definition at line 57 of file rules-meat.hpp.

7.36.3.3 get_seq()

Computes the sequence of free and locked cells in an BArray.

Parameters

а	An object of class BArray.
free	Pointer to a vector of pairs (i, j) listing the free cells.
locked	(optional) Pointer to a vector of pairs (i, j) listing the locked cells.

Returns

Nothing.

Definition at line 88 of file rules-meat.hpp.

7.36.3.4 operator()()

Check whether a given cell is free or locked.

Parameters

а	A BArray object
i	row position
j	col position

Returns

true If the cell is locked false If the cell is free

Definition at line 72 of file rules-meat.hpp.

7.36.3.5 operator=()

Definition at line 24 of file rules-meat.hpp.

7.36.3.6 size()

```
template<typename Array_Type , typename Data_Type >
uint Rules< Array_Type, Data_Type >::size ( ) const [inline], [noexcept]
```

Definition at line 75 of file rules-bones.hpp.

The documentation for this class was generated from the following files:

- include/barry/rules-bones.hpp
- include/barry/rules-meat.hpp

7.37 StatsCounter< Array_Type, Data_Type > Class Template Reference

Count stats for a single Array.

```
#include <statscounter-bones.hpp>
```

Public Member Functions

StatsCounter (const Array_Type *Array_)

Creator of a StatsCounter

StatsCounter (const StatsCounter< Array_Type, Data_Type > &counter)

Copy constructor.

• StatsCounter ()

Can be created without setting the array.

- ∼StatsCounter ()
- void reset_array (const Array_Type *Array_)

Changes the reference array for the counting.

- void add_counter (Counter< Array_Type, Data_Type > f_)
- void set_counters (Counters < Array_Type, Data_Type > *counters_)
- void count_init (uint i, uint j)

Counter functions This function recurses through the entries of Array and at each step of adding a new cell it uses the functions to list the statistics.

- void count current (uint i, uint i)
- std::vector< double > count_all ()
- Counters < Array_Type, Data_Type > * get_counters ()
- std::vector< std::string > get_names () const
- std::vector< std::string > get_descriptions () const
- size_t size () const

7.37.1 Detailed Description

```
template<typename Array_Type = BArray<>, typename Data_Type = bool> class StatsCounter< Array_Type, Data_Type >
```

Count stats for a single Array.

Users can a list of functions that can be used with this. The baseline set of arguments is a pointer to a binary array and a dataset to add the counts to.

Definition at line 19 of file statscounter-bones.hpp.

7.37.2 Constructor & Destructor Documentation

7.37.2.1 StatsCounter() [1/3]

Creator of a StatsCounter

Parameters

Array←	A const pointer to a BArray.

Definition at line 42 of file statscounter-bones.hpp.

7.37.2.2 StatsCounter() [2/3]

Copy constructor.

Parameters

counter

7.37.2.3 StatsCounter() [3/3]

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
StatsCounter< Array_Type, Data_Type >::StatsCounter ( ) [inline]
```

Can be created without setting the array.

Definition at line 64 of file statscounter-bones.hpp.

7.37.2.4 ~StatsCounter()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
StatsCounter< Array_Type, Data_Type >::~StatsCounter ( )
```

7.37.3 Member Function Documentation

7.37.3.1 add_counter()

7.37.3.2 count_all()

```
template<typename Array_Type , typename Data_Type >
std::vector< double > StatsCounter< Array_Type, Data_Type >::count_all [inline]
```

Definition at line 102 of file statscounter-meat.hpp.

7.37.3.3 count_current()

7.37.3.4 count_init()

Counter functions This function recurses through the entries of Array and at each step of adding a new cell it uses the functions to list the statistics.

7.37.3.5 get_counters()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Counters<Array_Type,Data_Type>* StatsCounter< Array_Type, Data_Type >::get_counters ( )
```

7.37.3.6 get_descriptions()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
std::vector< std::string > StatsCounter< Array_Type, Data_Type >::get_descriptions ( ) const
```

7.37.3.7 get_names()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
std::vector< std::string > StatsCounter< Array_Type, Data_Type >::get_names ( ) const
```

7.37.3.8 reset_array()

Changes the reference array for the counting.

Parameters

Array←	A pointer to an array of class Array_Type.
1_	

7.37.3.9 set_counters()

7.37.3.10 size()

```
template<typename Array_Type = BArray<>>, typename Data_Type = bool>
size_t StatsCounter< Array_Type, Data_Type >::size ( ) const [inline]
```

Definition at line 91 of file statscounter-bones.hpp.

The documentation for this class was generated from the following files:

- include/barry/statscounter-bones.hpp
- include/barry/statscounter-meat.hpp

7.38 Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data Rule Dyn Type > Class Template Reference

Compute the support of sufficient statistics.

```
#include <support-bones.hpp>
```

Public Member Functions

Support (const Array_Type &Array_)

Constructor passing a reference Array.

• Support (uint N_, uint M_)

Constructor specifying the dimensions of the array (empty).

- Support ()
- ∼Support ()
- void init_support (std::vector < Array_Type > *array_bank=nullptr, std::vector < double > *stats_bank=nullptr)
- void calc (std::vector< Array_Type > *array_bank=nullptr, std::vector< double > *stats_bank=nullptr, unsigned int max_num_elements_=0u)

Computes the entire support.

std::vector< double > get_counts () const

- std::vector< double > * get_current_stats () List current statistics.
- · void print () const
- const FreqTable & get data () const
- Counters < Array_Type, Data_Counter_Type > * get_counters ()

Vector of couter functions.

- Rules< Array_Type, Data_Rule_Type > * get_rules ()
- Vector of static rules (cells to iterate). Rules< Array_Type, Data_Rule_Dyn_Type > * get_rules_dyn ()

Vector of dynamic rules (to include/exclude a realizaton).

Resets the support calculator

If needed, the counters of a support object can be reused.

Parameters

Array↩	New array over which the support will be computed.	1
1_		

- void reset_array ()
- void reset_array (const Array_Type &Array_)

Manage counters

Parameters

f_	A counter to be added.
counters←	A vector of counters to be added.
_	

- void add_counter (Counter< Array_Type, Data_Counter_Type > f_)
- void set_counters (Counters < Array_Type, Data_Counter_Type > *counters_)

Manage rules

Parameters

f_	A rule to be added.
counters←	A vector of rules to be added.
_	

- void add_rule (Rule< Array_Type, Data_Rule_Type > *f_)
 void add_rule (Rule< Array_Type, Data_Rule_Type > f_)
 void set_rules (Rules< Array_Type, Data_Rule_Type > *rules_)
- void add_rule_dyn (Rule< Array_Type, Data_Rule_Dyn_Type > *f_)
 void add_rule_dyn (Rule< Array_Type, Data_Rule_Dyn_Type > f_)
- void set_rules_dyn (Rules < Array_Type, Data_Rule_Dyn_Type > *rules_)
- bool eval_rules_dyn (const std::vector< double > &counts, const uint &i, const uint &j)

Public Attributes

uint N

- uint M
- bool delete counters = true
- bool delete_rules = true
- bool delete_rules_dyn = true
- uint max num elements = BARRY MAX NUM ELEMENTS
- std::vector< double > current stats
- std::vector< size t > coordinates free
- std::vector< size_t > coordinates_locked
- size_t coordiantes_n_free
- size_t coordiantes_n_locked
- std::vector< double > change stats
- std::vector< size_t > hashes
- std::vector< bool > hashes initialized
- size_t n_counters

7.38.1 Detailed Description

```
template < typename Array_Type = BArray <>>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
class Support < Array_Type, Data_Counter_Type, Data_Rule_Dyn_Type >
```

Compute the support of sufficient statistics.

Given an array and a set of counters, this object iterates throughout the support set of the Array while at the same time computing the support of the sufficient statitics.

The members rule and rule_dyn allow constraining the support. The first will establish which cells of the array will be used to iterate, for example, in the case of social networks, self-loops are not allowed, so the entire diagonal would be fixed to zero, reducing the size of the support.

In the case of $rule_dyn$, the function will stablish dynamically whether the current state will be included in the counts or not. For example, this set of rules can be used to constrain the support to networks that have a prescribed degree sequence.

Definition at line 35 of file support-bones.hpp.

7.38.2 Constructor & Destructor Documentation

7.38.2.1 Support() [1/3]

Constructor passing a reference Array.

Definition at line 80 of file support-bones.hpp.

7.38.2.2 Support() [2/3]

Constructor specifying the dimensions of the array (empty).

Definition at line 89 of file support-bones.hpp.

7.38.2.3 Support() [3/3]

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::Support ( )
[inline]
```

Definition at line 96 of file support-bones.hpp.

7.38.2.4 ∼Support()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::~Support ()
[inline]
```

Definition at line 103 of file support-bones.hpp.

7.38.3 Member Function Documentation

7.38.3.1 add_counter()

7.38.3.2 add_rule() [1/2]

7.38.3.3 add_rule() [2/2]

7.38.3.4 add_rule_dyn() [1/2]

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarrow Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::add_rule_\leftarrow dyn ( Rule< Array_Type, Data_Rule_Dyn_Type > * f_- )
```

7.38.3.5 add_rule_dyn() [2/2]

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarrow Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::add_rule_\leftarrow dyn ( Rule< Array_Type, Data_Rule_Dyn_Type > f_- )
```

7.38.3.6 calc()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::calc (
    std::vector< Array_Type > * array_bank = nullptr,
    std::vector< double > * stats_bank = nullptr,
    unsigned int max_num_elements_ = 0u )
```

Computes the entire support.

Not to be used by the user. Sets the starting point in the array (column-major).

Parameters

array_bank	If specified, the counter will add to the vector each possible state of the array, as it counts.
stats_bank	If specified, the counter will add to the vector each possible set of statistics, as it counts.

7.38.3.7 eval_rules_dyn()

7.38.3.8 get_counters()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Counters<Array_Type, Data_Counter_Type>* Support< Array_Type, Data_Counter_Type, Data_Rule_←
Type, Data_Rule_Dyn_Type >::get_counters ()
```

Vector of couter functions.

7.38.3.9 get_counts()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< double > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn\times
_Type >::qet_counts ( ) const
```

7.38.3.10 get_current_stats()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< double >* Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_←
Dyn_Type >::get_current_stats ()
```

List current statistics.

7.38.3.11 get_data()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
const FreqTable& Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type
>::get_data ( ) const
```

7.38.3.12 get_rules()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Rules<Array_Type, Data_Rule_Type>* Support< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_rules ()
```

Vector of static rules (cells to iterate).

7.38.3.13 get_rules_dyn()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Rules<Array_Type, Data_Rule_Dyn_Type>* Support< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_rules_dyn ()
```

Vector of dynamic rules (to include/exclude a realizaton).

7.38.3.14 init support()

7.38.3.15 print()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::print ()
const
```

7.38.3.16 reset_array() [1/2]

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::reset_array
( )
```

7.38.3.17 reset_array() [2/2]

7.38.3.18 set counters()

7.38.3.19 set_rules()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::set_rules (
    Rules< Array_Type, Data_Rule_Type > * rules_ )
```

7.38.3.20 set_rules_dyn()

7.38.4 Member Data Documentation

7.38.4.1 change_stats

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< double > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn←
_Type >::change_stats
```

Definition at line 73 of file support-bones.hpp.

7.38.4.2 coordiantes_n_free

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::coordiantes
_n_free
```

Definition at line 71 of file support-bones.hpp.

7.38.4.3 coordiantes n locked

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::coordiantes\times
n locked
```

Definition at line 72 of file support-bones.hpp.

7.38.4.4 coordinates_free

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< size_t > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn←
_Type >::coordinates_free
```

Definition at line 69 of file support-bones.hpp.

7.38.4.5 coordinates_locked

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< size_t > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn←
_Type >::coordinates_locked
```

Definition at line 70 of file support-bones.hpp.

7.38.4.6 current stats

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< double > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn←
_Type >::current_stats
```

Definition at line 68 of file support-bones.hpp.

7.38.4.7 delete_counters

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_←
counters = true
```

Definition at line 62 of file support-bones.hpp.

7.38.4.8 delete rules

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_←
rules = true
```

Definition at line 63 of file support-bones.hpp.

7.38.4.9 delete_rules_dyn

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_←
rules_dyn = true
```

Definition at line 64 of file support-bones.hpp.

7.38.4.10 hashes

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< size_t > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn←
_Type >::hashes
```

Definition at line 74 of file support-bones.hpp.

7.38.4.11 hashes initialized

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< bool > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::hashes_initialized
```

Definition at line 75 of file support-bones.hpp.

7.38.4.12 M

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
uint Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::M
```

Definition at line 61 of file support-bones.hpp.

7.38.4.13 max_num_elements

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
uint Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::max_num_←
elements = BARRY_MAX_NUM_ELEMENTS
```

Definition at line 65 of file support-bones.hpp.

7.38.4.14 N

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
uint Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::N
```

Definition at line 61 of file support-bones.hpp.

7.38.4.15 n counters

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::n_←
counters
```

Definition at line 76 of file support-bones.hpp.

The documentation for this class was generated from the following file:

• include/barry/support-bones.hpp

7.39 vecHasher < T > Struct Template Reference

```
#include <typedefs.hpp>
```

Public Member Functions

• std::size_t operator() (std::vector< T > const &dat) const noexcept

7.39.1 Detailed Description

```
template < typename T> struct vecHasher < T>
```

Definition at line 106 of file typedefs.hpp.

7.39.2 Member Function Documentation

7.39.2.1 operator()()

Definition at line 109 of file typedefs.hpp.

The documentation for this struct was generated from the following file:

• include/barry/typedefs.hpp

Chapter 8

File Documentation

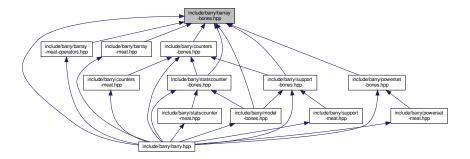
8.1 include/barry/barray-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "cell-bones.hpp"
#include "barraycell-bones.hpp"
Include dependency graph for barray-bones.hpp:
```



208 File Documentation

This graph shows which files directly or indirectly include this file:



Classes

class BArray < Cell_Type, Data_Type >
 Baseline class for binary arrays.

Macros

• #define BARRAY_BONES_HPP 1

8.1.1 Macro Definition Documentation

8.1.1.1 BARRAY_BONES_HPP

#define BARRAY_BONES_HPP 1

Definition at line 8 of file barray-bones.hpp.

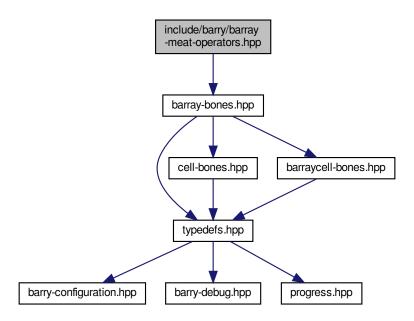
8.2 include/barry/barray-iterator.hpp File Reference

Classes

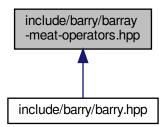
class ConstBArrayRowIter< Cell_Type, Data_Type >

8.3 include/barry/barray-meat-operators.hpp File Reference

#include "barray-bones.hpp"
Include dependency graph for barray-meat-operators.hpp:



This graph shows which files directly or indirectly include this file:



Macros

- #define BARRY_BARRAY_MEAT_OPERATORS_HPP 1
- #define BARRAY_TYPE() BArray<Cell_Type, Data_Type>
- #define BARRAY_TEMPLATE_ARGS() < typename Cell_Type, typename Data_Type>
- #define BARRAY_TEMPLATE(a, b) template BARRAY_TEMPLATE_ARGS() inline a BARRAY_TYPE()::b
- #define ROW(a) this->el_ij[a]
- #define COL(a) this->el_ji[a]

210 File Documentation

Functions

- template BARRAY_TEMPLATE_ARGS () inline void checkdim_(const BARRAY_TYPE() &lhs
- template const BARRAY_TYPE () &rhs)
- BARRAY_TEMPLATE (BARRAY_TYPE()&, operator+=)(const BArray< Cell_Type
- for (uint i=0u;i< nrow();++i) for(uint j=0u = el[POS(i, j)]
- j< ncol();++j) this-> operator() (i, j)+
- BARRAY_TEMPLATE (BARRAY_TYPE()&, operator+=)(const Cell_Type &rhs)
- BARRAY_TEMPLATE (BARRAY_TYPE()&, operator-=)(const BArray< Cell_Type
- BARRAY_TEMPLATE (BARRAY_TYPE()&, operator-=)(const Cell_Type &rhs)
- BARRAY TEMPLATE (BARRAY TYPE()&, operator*=)(const Cell Type &rhs)
- BARRAY_TEMPLATE (BARRAY_TYPE()&, operator/=)(const Cell_Type &rhs)

Variables

- Data_Type & rhs
- return * this

8.3.1 Macro Definition Documentation

8.3.1.1 BARRAY_TEMPLATE

Definition at line 11 of file barray-meat-operators.hpp.

8.3.1.2 BARRAY_TEMPLATE_ARGS

```
template BARRAY_TEMPLATE_ARGS() <typename Cell_Type, typename Data_Type>
```

Definition at line 9 of file barray-meat-operators.hpp.

8.3.1.3 BARRAY_TYPE

```
template Data_Type BARRAY_TYPE( ) BArray<Cell_Type, Data_Type>
```

Definition at line 7 of file barray-meat-operators.hpp.

8.3.1.4 BARRY_BARRAY_MEAT_OPERATORS_HPP

```
#define BARRY_BARRAY_MEAT_OPERATORS_HPP 1
```

Definition at line 5 of file barray-meat-operators.hpp.

8.3.1.5 COL

Definition at line 15 of file barray-meat-operators.hpp.

8.3.1.6 ROW

Definition at line 14 of file barray-meat-operators.hpp.

8.3.2 Function Documentation

8.3.2.1 BARRAY_TEMPLATE() [1/6]

Definition at line 88 of file barray-meat-operators.hpp.

8.3.2.2 BARRAY_TEMPLATE() [2/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator+ ) const
```

212 File Documentation

8.3.2.3 BARRAY_TEMPLATE() [3/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator+ ) const &
```

Definition at line 46 of file barray-meat-operators.hpp.

8.3.2.4 BARRAY_TEMPLATE() [4/6]

8.3.2.5 BARRAY_TEMPLATE() [5/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator- ) const &
```

Definition at line 75 of file barray-meat-operators.hpp.

8.3.2.6 BARRAY_TEMPLATE() [6/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator/ ) const &
```

Definition at line 105 of file barray-meat-operators.hpp.

8.3.2.7 BARRAY_TEMPLATE_ARGS()

```
template BARRAY_TEMPLATE_ARGS ( ) const &
```

8.3.2.8 BARRAY_TYPE()

```
template const BARRAY_TYPE ( ) &
```

Definition at line 20 of file barray-meat-operators.hpp.

8.3.2.9 for()

```
for ( ) = el[POS(i, j)] [pure virtual]
```

Definition at line 66 of file barray-meat-operators.hpp.

8.3.2.10 operator()()

8.3.3 Variable Documentation

8.3.3.1 rhs

```
Data_Type & rhs
Initial value:
{
    checkdim_(*this, rhs)
```

Definition at line 33 of file barray-meat-operators.hpp.

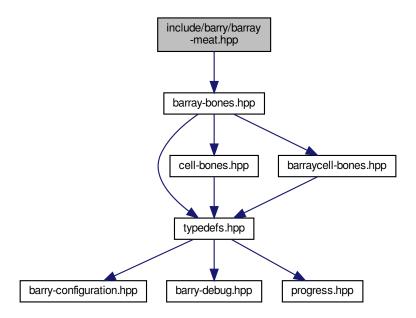
8.3.3.2 this

```
return * this
```

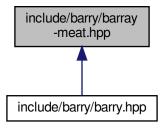
Definition at line 43 of file barray-meat-operators.hpp.

8.4 include/barry/barray-meat.hpp File Reference

#include "barray-bones.hpp"
Include dependency graph for barray-meat.hpp:



This graph shows which files directly or indirectly include this file:



Macros

- #define BARRAY_TYPE() BArray<Cell_Type, Data_Type>
- #define BARRAY_TEMPLATE_ARGS() < typename Cell_Type, typename Data_Type>
- #define BARRAY_TEMPLATE(a, b) template BARRAY_TEMPLATE_ARGS() inline a BARRAY_TYPE()::b
- #define ROW(a) this->el_ij[a]
- #define COL(a) this->el_ji[a]

Functions

```
• BARRAY TEMPLATE (, BArray)(uint N_
• el ij resize (N)
• el_ji resize (M)

    for (uint i=0u;i< source.size();++i)</li>

    Data Type bool M (Array .M)

    BARRAY_TEMPLATE (BARRAY_TYPE() &, operator=)(const BArray< Cell_Type</li>

    BARRAY_TEMPLATE (, BArray)(BARRAY_TYPE() &&x) noexcept

    BARRAY TEMPLATE (BARRAY TYPE() &, operator=)(BARRAY TYPE() &&x) noexcept

• BARRAY TEMPLATE (bool, operator==)(const BARRAY TYPE() & Array )

    BARRAY TEMPLATE (,~BArray)()

    BARRAY_TEMPLATE (void, set_data)(Data_Type *data_

    BARRAY TEMPLATE (Data Type *, D ptr)()

• BARRAY_TEMPLATE (Data_Type &, D)()

    BARRAY TEMPLATE (void, out of range)(uint i

    BARRAY TEMPLATE (Cell Type, get cell)(uint i

    if (ROW(i).size()==0u) return(Cell_Type) 0.0

• if (search !=ROW(i).end()) return search -> second.value
• return (Cell_Type) 0.0

    BARRAY_TEMPLATE (std::vector< Cell_Type >, get_row_vec)(uint i

    std::vector< Cell Type > ans (ncol(),(Cell Type) false)

    for (const auto &iter :row(i, false)) ans[iter.first]

    BARRAY_TEMPLATE (void, get_row_vec)(std

    BARRAY_TEMPLATE (BARRAY_TYPE() &, operator-=)(const std

• BARRAY_TEMPLATE (void, insert_cell)(uint i
· if (check exists)

    COL (j).emplace(i

• & ROW (i)[j])

    BARRAY_TEMPLATE (void, swap_cells)(uint i0

if (report !=nullptr)(*report)
• if (check0 &check1)

    else if (!check0 &check1)

• else if (check0 &!check1)

    BARRAY TEMPLATE (void, toggle cell)(uint i

• BARRAY_TEMPLATE (void, swap_rows)(uint i0
if (ROW(i0).size()==0u) move0
if (ROW(i1).size()==0u) move1
• if (!move0 &&!move1) return

    ROW (i0).swap(ROW(i1))

    BARRAY_TEMPLATE (void, swap_cols)(uint j0

• if (COL(j0).size()==0u) check0
if (COL(j1).size()==0u) check1

    if (check0 &&check1)

    else if (check0 &&!check1)

    else if (!check0 &&check1)

• BARRAY_TEMPLATE (void, zero_row)(uint i

    for (auto row=row0.begin();row !=row0.end();++row) rm cell(i

    BARRAY_TEMPLATE (void, zero_col)(uint j

    if (COL(j).size()==0u) return

• BARRAY_TEMPLATE (void, transpose)()

    BARRAY_TEMPLATE (void, clear)(bool hard)

    BARRAY_TEMPLATE (void, resize)(uint N_

• if (M < M) for (uint j = N)
```

Variables

```
    uint M
```

- uint const std::vector< uint > & source
- uint const std::vector< uint > const std::vector< uint > & target
- uint const std::vector< uint > const std::vector< cell_Type > & value
- uint const std::vector< uint > const std::vector< Cell Type > bool add
- if(source.size() !=value.size()) throw std N = N_
- M = M
- return
- Data_Type & Array_
- Data_Type bool copy_data
- bool delete_data_
- data = data
- delete_data = delete_data_
- uint j const
- uint j
- auto search = ROW(i).find(j)
- · return ans
- uint const Cell
 Cell_Type > & v
- uint const Cell< Cell_Type > bool check_bounds
- uint const Cell
 Cell_Type > bool bool check_exists
- else
- NCells
- uint j0
- uint uint i1
- uint uint uint j1
- uint uint bool int int * report
- auto row0 = ROW(i)
- row first
- · row false
- auto col0 = COL(j)

8.4.1 Macro Definition Documentation

8.4.1.1 BARRAY_TEMPLATE

Definition at line 17 of file barray-meat.hpp.

8.4.1.2 BARRAY_TEMPLATE_ARGS

```
#define BARRAY_TEMPLATE_ARGS( ) <typename Cell_Type, typename Data_Type>
```

Definition at line 15 of file barray-meat.hpp.

8.4.1.3 BARRAY_TYPE

```
#define BARRAY_TYPE( ) BArray<Cell_Type, Data_Type>
```

Definition at line 13 of file barray-meat.hpp.

8.4.1.4 COL

Definition at line 21 of file barray-meat.hpp.

8.4.1.5 ROW

Definition at line 20 of file barray-meat.hpp.

8.4.2 Function Documentation

8.4.2.1 ans()

8.4.2.2 BARRAY_TEMPLATE() [1/24]

```
BARRAY_TEMPLATE (

BArray ) && [noexcept]
```

Definition at line 230 of file barray-meat.hpp.

8.4.2.3 BARRAY_TEMPLATE() [2/24]

```
BARRAY_TEMPLATE (
BArray )
```

8.4.2.4 BARRAY_TEMPLATE() [3/24]

```
BARRAY_TEMPLATE ( \sim \textit{BArray} \ )
```

Definition at line 339 of file barray-meat.hpp.

8.4.2.5 BARRAY_TEMPLATE() [4/24]

Definition at line 597 of file barray-meat.hpp.

8.4.2.6 BARRAY_TEMPLATE() [5/24]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE() & ,
          operator ) && [noexcept]
```

Definition at line 272 of file barray-meat.hpp.

8.4.2.7 BARRAY_TEMPLATE() [6/24]

8.4.2.8 BARRAY_TEMPLATE() [7/24]

```
BARRAY_TEMPLATE (
          bool ,
          operator = = ) const &
```

Definition at line 321 of file barray-meat.hpp.

8.4.2.9 BARRAY_TEMPLATE() [8/24]

8.4.2.10 BARRAY_TEMPLATE() [9/24]

```
BARRAY_TEMPLATE (

Data_Type & ,

D )
```

Definition at line 372 of file barray-meat.hpp.

8.4.2.11 BARRAY_TEMPLATE() [10/24]

Definition at line 361 of file barray-meat.hpp.

8.4.2.12 BARRAY_TEMPLATE() [11/24]

```
BARRAY_TEMPLATE (
          std::vector< Cell_Type > ,
          get_row_vec )
```

8.4.2.13 BARRAY_TEMPLATE() [12/24]

```
BARRAY_TEMPLATE ( void , clear )
```

Definition at line 1130 of file barray-meat.hpp.

8.4.2.14 BARRAY_TEMPLATE() [13/24]

Definition at line 452 of file barray-meat.hpp.

8.4.2.15 BARRAY_TEMPLATE() [14/24]

8.4.2.16 BARRAY_TEMPLATE() [15/24]

8.4.2.17 BARRAY_TEMPLATE() [16/24]

8.4.2.18 BARRAY_TEMPLATE() [17/24]

8.4.2.19 BARRAY_TEMPLATE() [18/24]

8.4.2.20 BARRAY_TEMPLATE() [19/24]

8.4.2.21 BARRAY_TEMPLATE() [20/24]

8.4.2.22 BARRAY_TEMPLATE() [21/24]

8.4.2.23 BARRAY_TEMPLATE() [22/24]

Definition at line 1069 of file barray-meat.hpp.

8.4.2.24 BARRAY_TEMPLATE() [23/24]

```
BARRAY_TEMPLATE ( void , zero_col )
```

8.4.2.25 BARRAY_TEMPLATE() [24/24]

```
BARRAY_TEMPLATE (

void ,

zero_row )
```

```
8.4.2.26 COL()
```

```
COL (
```

8.4.2.27 for() [1/3]

```
for (
    auto row = row0.begin();row !=row0.end();++row )
```

8.4.2.28 for() [2/3]

8.4.2.29 for() [3/3]

```
for ( )
```

Definition at line 51 of file barray-meat.hpp.

8.4.2.30 if() [1/17]

```
else if ( !check0 && check1 )
```

Definition at line 1008 of file barray-meat.hpp.

8.4.2.31 if() [2/17]

```
else if (
    !check0 & check1 )
```

Definition at line 856 of file barray-meat.hpp.

8.4.2.32 if() [3/17]

```
if (
    !move0 &&! move1 )
```

8.4.2.33 if() [4/17]

Definition at line 864 of file barray-meat.hpp.

8.4.2.34 if() [5/17]

```
else if (
          check0 &&! check1 )
```

Definition at line 999 of file barray-meat.hpp.

8.4.2.35 if() [6/17]

```
if ( check0 && check1)
```

Definition at line 972 of file barray-meat.hpp.

8.4.2.36 if() [7/17]

```
if ( check0 & check1)
```

Definition at line 838 of file barray-meat.hpp.

8.4.2.37 if() [8/17]

Definition at line 679 of file barray-meat.hpp.

```
8.4.2.38 if() [9/17]
```

```
if ( COL(j).size() = =0u )
```

8.4.2.39 if() [10/17]

```
if ( \label{eq:col} \mathtt{COL}(\mathtt{j0}).\mathtt{size}() \ = = 0u \ )
```

8.4.2.40 if() [11/17]

```
if ( \label{eq:col} \mathtt{COL(j1).size()} \ = \ = 0u \ )
```

8.4.2.41 if() [12/17]

```
else if ( ) = N_
```

Definition at line 86 of file barray-meat.hpp.

8.4.2.42 if() [13/17]

```
if (  {\tt report !} \quad = {\tt nullptr} \; ) \\
```

8.4.2.43 if() [14/17]

```
if ( \label{eq:row_row_row} \mbox{ROW(i).size()} \ = \ = \mbox{$0$$u$} \ )
```

8.4.2.44 if() [15/17]

```
if ( \label{eq:row_row_row} \text{ROW(iO).size()} \quad = = 0u \text{ )}
```

8.4.2.45 if() [16/17]

```
if ( \label{eq:row_row_row} \text{ROW(i1).size()} \quad = = 0u \text{ )}
```

8.4.2.46 if() [17/17]

```
if (
    search ! = ROW(i).end() ) -> second.value
```

8.4.2.47 M()

```
Data_Type bool M ( \label{eq:continuous} \text{Array}.\quad \textit{M}\ )
```

Definition at line 136 of file barray-meat.hpp.

8.4.2.48 resize() [1/2]

```
el_ji resize (
M )
```

8.4.2.49 resize() [2/2]

```
el_ij resize (
N )
```

8.4.2.50 return()

8.4.2.51 ROW() [1/2]

```
& ROW ( i )
```

8.4.2.52 ROW() [2/2]

```
ROW ( i0 )
```

8.4.3 Variable Documentation

8.4.3.1 add

```
uint const std::vector< uint > const std::vector< uint > bool add

Initial value:
{
    if (source.size() != target.size())
        throw std::length_error("-source- and -target- don't match on length.")
```

Definition at line 34 of file barray-meat.hpp.

8.4.3.2 ans

return ans

Definition at line 449 of file barray-meat.hpp.

8.4.3.3 Array_

```
Data_Type & Array_
```

Definition at line 134 of file barray-meat.hpp.

8.4.3.4 check_bounds

```
bool check_bounds

Initial value:
{
    if (check_bounds) {
        out_of_range(i0,0u);
        out_of_range(i1,0u);
    }
    bool move0=true, move1=true
```

Definition at line 672 of file barray-meat.hpp.

8.4.3.5 check_exists

```
uint bool int check_exists
Initial value:
{
    if (check_bounds)
        out_of_range(i,j)
```

Definition at line 673 of file barray-meat.hpp.

8.4.3.6 col0

```
auto col0 = COL(j)
```

Definition at line 1061 of file barray-meat.hpp.

8.4.3.7 const

```
uint bool check_bounds const

Initial value:
{
    if (i >= N)
        throw std::range_error("The row is out of range.")
```

Definition at line 402 of file barray-meat.hpp.

8.4.3.8 copy_data

```
Data_Type bool copy_data
```

Definition at line 135 of file barray-meat.hpp.

8.4.3.9 data

```
data = data_
```

Definition at line 354 of file barray-meat.hpp.

8.4.3.10 delete_data

```
delete_data = delete_data_
```

Definition at line 355 of file barray-meat.hpp.

8.4.3.11 delete_data_

Definition at line 348 of file barray-meat.hpp.

8.4.3.12 else

Definition at line 703 of file barray-meat.hpp.

8.4.3.13 false

```
row false
```

Definition at line 1042 of file barray-meat.hpp.

8.4.3.14 first

```
row first
```

Definition at line 1042 of file barray-meat.hpp.

8.4.3.15 i1

```
uint i1
```

Definition at line 776 of file barray-meat.hpp.

8.4.3.16 j

```
uint j
```

Initial value:

```
if (init_fun == nullptr)
    return 0.0
```

Definition at line 414 of file barray-meat.hpp.

8.4.3.17 j0

```
uint j0
```

Definition at line 775 of file barray-meat.hpp.

8.4.3.18 j1

```
uint j1
```

Definition at line 776 of file barray-meat.hpp.

8.4.3.19 M

```
M = M_{\underline{}}
```

Definition at line 44 of file barray-meat.hpp.

8.4.3.20 M_

```
uint M_
```

Initial value:

{

```
if (N_ < N)
    for (uint i = N_; i < N; ++i)
        zero_row(i, false)</pre>
```

Definition at line 30 of file barray-meat.hpp.

8.4.3.21 N

```
if (source.size() != target.size()) throw std if (source.size() != value.size()) throw std N = N
```

Definition at line 43 of file barray-meat.hpp.

8.4.3.22 NCells

NCells

Definition at line 707 of file barray-meat.hpp.

8.4.3.23 report

```
uint uint uint bool int int* report
```

Definition at line 779 of file barray-meat.hpp.

8.4.3.24 return

return

Definition at line 66 of file barray-meat.hpp.

8.4.3.25 row0

```
auto row0 = ROW(i)
```

Definition at line 1040 of file barray-meat.hpp.

8.4.3.26 search

```
auto search = ROW(i).find(j)
```

Definition at line 426 of file barray-meat.hpp.

8.4.3.27 source

```
uint const std::vector< uint > & source
```

Definition at line 31 of file barray-meat.hpp.

8.4.3.28 target

```
uint const std::vector< uint > const std::vector< uint > & target
```

Definition at line 32 of file barray-meat.hpp.

8.4.3.29 v

```
uint Cell_Type v
```

Definition at line 671 of file barray-meat.hpp.

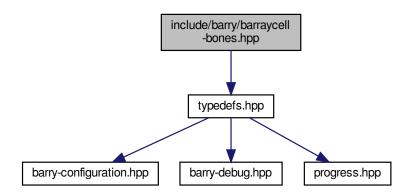
8.4.3.30 value

uint const std::vector< uint > const std::vector< uint > const std::vector< Cell_Type >& value

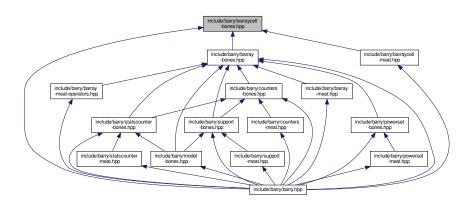
Definition at line 33 of file barray-meat.hpp.

8.5 include/barry/barraycell-bones.hpp File Reference

#include "typedefs.hpp"
Include dependency graph for barraycell-bones.hpp:



This graph shows which files directly or indirectly include this file:

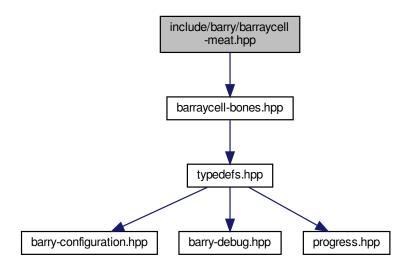


Classes

- class BArrayCell
 Cell_Type, Data_Type
- class BArrayCell_const< Cell_Type, Data_Type >

8.6 include/barry/barraycell-meat.hpp File Reference

#include "barraycell-bones.hpp"
Include dependency graph for barraycell-meat.hpp:



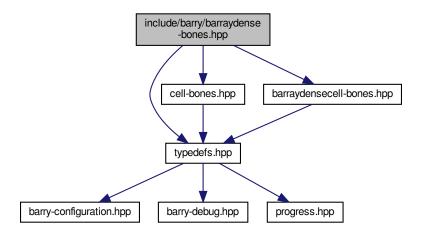
This graph shows which files directly or indirectly include this file:



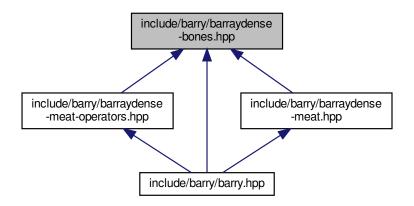
8.7 include/barry/barraydense-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "cell-bones.hpp"
```

#include "barraydensecell-bones.hpp"
Include dependency graph for barraydense-bones.hpp:



This graph shows which files directly or indirectly include this file:



Classes

class BArrayDense < Cell_Type, Data_Type >
 Baseline class for binary arrays.

Macros

• #define BARRY_BARRAYDENSE_BONES_HPP 1

8.7.1 Macro Definition Documentation

8.7.1.1 BARRY_BARRAYDENSE_BONES_HPP

#define BARRY_BARRAYDENSE_BONES_HPP 1

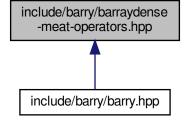
Definition at line 8 of file barraydense-bones.hpp.

8.8 include/barry/barraydense-meat-operators.hpp File Reference

#include "barraydense-bones.hpp"
Include dependency graph for barraydense-meat-operators.hpp:



This graph shows which files directly or indirectly include this file:



Macros

- #define BARRY_BARRAYDENSE_MEAT_OPERATORS_HPP 1
- #define BDENSE_TYPE() BArrayDense<Cell_Type, Data_Type>
- #define BDENSE_TEMPLATE_ARGS() < typename Cell_Type, typename Data_Type>
- #define BDENSE TEMPLATE(a, b) template BDENSE TEMPLATE ARGS() inline a BDENSE TYPE()::b
- #define ROW(a) this->el ij[a]
- #define COL(a) this->el_ji[a]
- #define POS(a, b) (b)*N + (a)
- #define POS_N(a, b, c) (b)*(c) + (a)

Functions

- template BDENSE_TEMPLATE_ARGS () inline void checkdim_(const BDENSE_TYPE() &lhs
- template const BDENSE_TYPE () &rhs)
- BDENSE TEMPLATE (BDENSE TYPE()&, operator+=)(const BDENSE TYPE() &rhs)
- BDENSE_TEMPLATE (BDENSE_TYPE()&, operator-=)(const BDENSE_TYPE() &rhs)
- BDENSE_TEMPLATE (BDENSE_TYPE()&, operator*=)(const Cell_Type &rhs)
- BDENSE_TEMPLATE (BDENSE_TYPE()&, operator/=)(const Cell_Type &rhs)

8.8.1 Macro Definition Documentation

8.8.1.1 BARRY_BARRAYDENSE_MEAT_OPERATORS_HPP

```
#define BARRY_BARRAYDENSE_MEAT_OPERATORS_HPP 1
```

Definition at line 5 of file barraydense-meat-operators.hpp.

8.8.1.2 BDENSE_TEMPLATE

Definition at line 11 of file barraydense-meat-operators.hpp.

8.8.1.3 BDENSE_TEMPLATE_ARGS

```
template BDENSE_TEMPLATE_ARGS() <typename Cell_Type, typename Data_Type>
```

Definition at line 9 of file barraydense-meat-operators.hpp.

8.8.1.4 BDENSE_TYPE

```
template Data_Type BDENSE_TYPE() BArrayDense<Cell_Type, Data_Type>
```

Definition at line 7 of file barraydense-meat-operators.hpp.

8.8.1.5 COL

Definition at line 15 of file barraydense-meat-operators.hpp.

8.8.1.6 POS

```
#define POS( \label{eq:a_b} a, \\ b \ ) \ (b)*N + (a)
```

Definition at line 16 of file barraydense-meat-operators.hpp.

8.8.1.7 POS N

Definition at line 17 of file barraydense-meat-operators.hpp.

8.8.1.8 ROW

Definition at line 14 of file barraydense-meat-operators.hpp.

8.8.2 Function Documentation

8.8.2.1 BDENSE_TEMPLATE() [1/4]

Definition at line 90 of file barraydense-meat-operators.hpp.

8.8.2.2 BDENSE_TEMPLATE() [2/4]

Definition at line 34 of file barraydense-meat-operators.hpp.

8.8.2.3 BDENSE_TEMPLATE() [3/4]

Definition at line 61 of file barraydense-meat-operators.hpp.

8.8.2.4 BDENSE_TEMPLATE() [4/4]

Definition at line 101 of file barraydense-meat-operators.hpp.

8.8.2.5 BDENSE_TEMPLATE_ARGS()

```
template BDENSE_TEMPLATE_ARGS ( ) const &
```

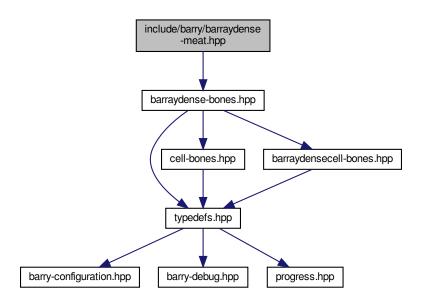
8.8.2.6 BDENSE_TYPE()

```
template const BDENSE_TYPE ( ) &
```

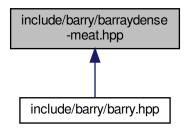
Definition at line 22 of file barraydense-meat-operators.hpp.

8.9 include/barry/barraydense-meat.hpp File Reference

#include "barraydense-bones.hpp"
Include dependency graph for barraydense-meat.hpp:



This graph shows which files directly or indirectly include this file:



Macros

- #define BDENSE_TYPE() BArrayDense<Cell_Type, Data_Type>
- #define BDENSE_TEMPLATE_ARGS() < typename Cell_Type, typename Data_Type>
- #define BDENSE_TEMPLATE(a, b) template BDENSE_TEMPLATE_ARGS() inline a BDENSE_TYPE()::b
- #define ROW(a) this->el_ij[a]
- #define COL(a) this->el_ji[a]
- #define POS(a, b) (b)*N + (a)
- #define POS_N(a, b, c) (b)*(c) + (a)
- #define ZERO_CELL static_cast<Cell_Type>(0.0)

Functions

```
    BDENSE_TEMPLATE (, BArrayDense)(uint N_

    el resize (N *M, ZERO_CELL)

• el_rowsums resize (N, ZERO_CELL)
• el colsums resize (M, ZERO CELL)

    for (uint i=0u;i< source.size();++i)</li>

    BDENSE TEMPLATE (, BArrayDense)(const BDENSE TYPE() & Array

• bool M (Array .M)
• BDENSE_TEMPLATE (BDENSE_TYPE() &, operator=)(const BDENSE_TYPE() &Array_)
• BDENSE TEMPLATE (, BArrayDense)(BDENSE TYPE() &&x) noexcept

    BDENSE TEMPLATE (BDENSE TYPE() &, operator=)(BDENSE TYPE() &&x) noexcept

• BDENSE_TEMPLATE (bool, operator==)(const BDENSE_TYPE() &Array_)

    BDENSE TEMPLATE (, ~BArrayDense)()

    BDENSE_TEMPLATE (void, set_data)(Data_Type *data_

• BDENSE_TEMPLATE (Data_Type *, D_ptr)()
• BDENSE TEMPLATE (const Data Type *, D ptr)() const
• BDENSE TEMPLATE (Data Type &, D)()

    BDENSE TEMPLATE (const Data Type &, D)() const

• BDENSE_TEMPLATE (void, out_of_range)(uint i

    BDENSE_TEMPLATE (Cell_Type, get_cell)(uint i

• BDENSE_TEMPLATE (std::vector< Cell_Type >, get_row_vec)(uint i
• std::vector< Cell Type > ans (ncol(), static cast< Cell Type > (false))
• BDENSE TEMPLATE (void, get row vec)(std

    BDENSE_TEMPLATE (Entries < Cell_Type >, get_entries)() const

    BDENSE_TEMPLATE (bool, is_empty)(uint i

    BDENSE_TEMPLATE (unsigned int, nrow)() const noexcept

• BDENSE TEMPLATE (unsigned int, ncol)() const noexcept
• BDENSE TEMPLATE (unsigned int, nnozero)() const noexcept
• BDENSE TEMPLATE (Cell< Cell Type >, default val)() const
• BDENSE TEMPLATE (BDENSE TYPE() &, operator+=)(const std

    BDENSE TEMPLATE (BDENSE TYPE() &, operator-=)(const std

    BDENSE_TEMPLATE (void, insert_cell)(uint i

• if (el[POS(i, j)]==BARRY_ZERO_DENSE)

    BDENSE TEMPLATE (void, swap cells)(uint i0

• if ((i0==i1) &&(j0==j1)) return
• rm cell (i0, j0, false, false)
• rm cell (i1, j1, false, false)
• insert_cell (i0, j0, val1, false, false)

    insert cell (i1, j1, val0, false, false)

• BDENSE TEMPLATE (void, toggle cell)(uint i
· else rm cell (i, j, false, false)

    BDENSE_TEMPLATE (void, swap_rows)(uint i0

    BDENSE TEMPLATE (void, swap cols)(uint j0

• BDENSE_TEMPLATE (void, zero_row)(uint i
• if (el rowsums[i]==ZERO CELL) return

    BDENSE TEMPLATE (void, zero col)(uint j

• if (el_colsums[j]==ZERO_CELL) return

    BDENSE TEMPLATE (void, transpose)()

    BDENSE_TEMPLATE (void, clear)(bool hard)

• BDENSE TEMPLATE (void, resize)(uint N
• el resize (N *M , ZERO CELL)
• el rowsums resize (N, ZERO CELL)
```

el_colsums resize (M_, ZERO_CELL)BDENSE_TEMPLATE (void, reserve)()

```
• BDENSE_TEMPLATE (void, print)(const char *fmt

    va_start (args, fmt)

    vprintf (fmt, args)

    va_end (args)
    • BDENSE TEMPLATE (const std::vector< Cell Type > &, get data)() const
    • BDENSE_TEMPLATE (const Cell_Type, rowsum)(unsigned int i) const
   • BDENSE TEMPLATE (const Cell Type, colsum)(unsigned int j) const
Variables

    uint M

    • uint const std::vector< uint > & source

    uint const std::vector< uint > const std::vector< uint > & target

    uint const std::vector< uint > const std::vector< uint > const std::vector< Cell Type > & value

    uint const std::vector< uint > const std::vector< Cell_Type > bool add

    if(source.size() !=value.size()) throw std N = N

    • M = M_

    return

   · bool copy_data
    · bool delete_data_
    • data = data_

    delete data = delete data

   · uint j const

    uint j

    return el [POS(i, j)] == ZERO_CELL
    · return ans
   • uint const Cell< Cell_Type > & v

    uint const Cell
    Cell_Type > bool check_bounds

    uint const Cell< Cell_Type > bool bool check_exists

    else

    • el rowsums [i] = (v.value - old)
    • el_colsums [j] = (v.value - old)
   • uint j0
   • uint uint i1
   · uint uint uint j1
```

8.9.1 Macro Definition Documentation

uint uint uint bool int int * report
Cell_Type val0 = el[POS(i0,j0)]
Cell_Type val1 = el[POS(i1,j1)]

8.9.1.1 BDENSE TEMPLATE

Definition at line 27 of file barraydense-meat.hpp.

falsecol

8.9.1.2 BDENSE_TEMPLATE_ARGS

```
#define BDENSE_TEMPLATE_ARGS( ) <typename Cell_Type, typename Data_Type>
```

Definition at line 25 of file barraydense-meat.hpp.

8.9.1.3 BDENSE_TYPE

```
#define BDENSE_TYPE( ) BArrayDense<Cell_Type, Data_Type>
```

Definition at line 23 of file barraydense-meat.hpp.

8.9.1.4 COL

Definition at line 31 of file barraydense-meat.hpp.

8.9.1.5 POS

```
#define POS(  a, \\ b ) \ (b)*N + (a)
```

Definition at line 32 of file barraydense-meat.hpp.

8.9.1.6 POS_N

Definition at line 33 of file barraydense-meat.hpp.

8.9.1.7 ROW

Definition at line 30 of file barraydense-meat.hpp.

8.9.1.8 ZERO_CELL

```
#define ZERO_CELL static_cast<Cell_Type>(0.0)
```

Definition at line 38 of file barraydense-meat.hpp.

8.9.2 Function Documentation

8.9.2.1 ans()

8.9.2.2 BDENSE_TEMPLATE() [1/39]

Definition at line 240 of file barraydense-meat.hpp.

8.9.2.3 BDENSE_TEMPLATE() [2/39]

8.9.2.4 BDENSE_TEMPLATE() [3/39]

```
BDENSE_TEMPLATE (
BArrayDense )
```

8.9.2.5 BDENSE_TEMPLATE() [4/39]

```
BDENSE_TEMPLATE ( \sim BArrayDense )
```

Definition at line 318 of file barraydense-meat.hpp.

8.9.2.6 BDENSE_TEMPLATE() [5/39]

Definition at line 566 of file barraydense-meat.hpp.

8.9.2.7 BDENSE_TEMPLATE() [6/39]

Definition at line 584 of file barraydense-meat.hpp.

8.9.2.8 BDENSE_TEMPLATE() [7/39]

Definition at line 257 of file barraydense-meat.hpp.

8.9.2.9 BDENSE_TEMPLATE() [8/39]

Definition at line 194 of file barraydense-meat.hpp.

8.9.2.10 BDENSE_TEMPLATE() [9/39]

```
BDENSE_TEMPLATE (
          bool ,
          is_empty )
```

8.9.2.11 BDENSE_TEMPLATE() [10/39]

```
BDENSE_TEMPLATE (
          bool ,
          operator = = ) const &
```

Definition at line 300 of file barraydense-meat.hpp.

8.9.2.12 BDENSE_TEMPLATE() [11/39]

Definition at line 562 of file barraydense-meat.hpp.

8.9.2.13 BDENSE_TEMPLATE() [12/39]

8.9.2.14 BDENSE_TEMPLATE() [13/39]

Definition at line 999 of file barraydense-meat.hpp.

8.9.2.15 BDENSE_TEMPLATE() [14/39]

Definition at line 994 of file barraydense-meat.hpp.

8.9.2.16 BDENSE_TEMPLATE() [15/39]

Definition at line 353 of file barraydense-meat.hpp.

8.9.2.17 BDENSE_TEMPLATE() [16/39]

Definition at line 345 of file barraydense-meat.hpp.

8.9.2.18 BDENSE_TEMPLATE() [17/39]

Definition at line 989 of file barraydense-meat.hpp.

8.9.2.19 BDENSE_TEMPLATE() [18/39]

Definition at line 349 of file barraydense-meat.hpp.

8.9.2.20 BDENSE_TEMPLATE() [19/39]

Definition at line 341 of file barraydense-meat.hpp.

8.9.2.21 BDENSE_TEMPLATE() [20/39]

Definition at line 502 of file barraydense-meat.hpp.

8.9.2.22 BDENSE_TEMPLATE() [21/39]

8.9.2.23 BDENSE_TEMPLATE() [22/39]

```
BDENSE_TEMPLATE (
         unsigned int ,
         ncol ) const [noexcept]
```

Definition at line 548 of file barraydense-meat.hpp.

8.9.2.24 BDENSE_TEMPLATE() [23/39]

```
BDENSE_TEMPLATE (
          unsigned int ,
          nnozero ) const [noexcept]
```

Definition at line 552 of file barraydense-meat.hpp.

8.9.2.25 BDENSE_TEMPLATE() [24/39]

```
BDENSE_TEMPLATE (
         unsigned int ,
         nrow ) const [noexcept]
```

Definition at line 544 of file barraydense-meat.hpp.

8.9.2.26 BDENSE_TEMPLATE() [25/39]

```
BDENSE_TEMPLATE (
     void ,
     clear )
```

Definition at line 896 of file barraydense-meat.hpp.

8.9.2.27 BDENSE_TEMPLATE() [26/39]

Definition at line 402 of file barraydense-meat.hpp.

8.9.2.28 BDENSE_TEMPLATE() [27/39]

8.9.2.29 BDENSE_TEMPLATE() [28/39]

8.9.2.30 BDENSE_TEMPLATE() [29/39]

8.9.2.31 BDENSE_TEMPLATE() [30/39]

```
BDENSE_TEMPLATE (
     void ,
     reserve )
```

Definition at line 946 of file barraydense-meat.hpp.

8.9.2.32 BDENSE_TEMPLATE() [31/39]

```
BDENSE_TEMPLATE (
     void ,
     resize )
```

8.9.2.33 BDENSE_TEMPLATE() [32/39]

8.9.2.34 BDENSE_TEMPLATE() [33/39]

8.9.2.35 BDENSE_TEMPLATE() [34/39]

8.9.2.36 BDENSE_TEMPLATE() [35/39]

8.9.2.37 BDENSE_TEMPLATE() [36/39]

8.9.2.38 BDENSE_TEMPLATE() [37/39]

```
BDENSE_TEMPLATE (
            void ,
            transpose )
```

Definition at line 868 of file barraydense-meat.hpp.

8.9.2.39 BDENSE_TEMPLATE() [38/39]

8.9.2.40 BDENSE_TEMPLATE() [39/39]

8.9.2.41 for()

```
for ( )
```

Definition at line 64 of file barraydense-meat.hpp.

```
8.9.2.42 if() [1/4]
```

```
if (  ( {\tt i0 == i1}) \ \&\& \, ( {\tt j0 == j1}) \ )
```

8.9.2.43 if() [2/4]

```
if (  \mbox{el } [POS(i,\ j)] \ = \ = \ BARRY\_ZERO\_DENSE \ )
```

Definition at line 663 of file barraydense-meat.hpp.

8.9.2.44 if() [3/4]

```
if ( {\tt el\_colsums}~[j] = {\tt =ZERO\_CELL}~)
```

8.9.2.45 if() [4/4]

```
if (
    el_rowsums [i] = =ZERO_CELL )
```

8.9.2.46 insert_cell() [1/2]

8.9.2.47 insert_cell() [2/2]

8.9.2.48 M()

```
bool M ( \label{eq:Array_.} \mbox{Array}. \mbox{$M$} \mbox{)}
```

Definition at line 157 of file barraydense-meat.hpp.

8.9.2.49 resize() [1/6]

8.9.2.50 resize() [2/6]

8.9.2.51 resize() [3/6]

```
el resize ( \label{eq:N*M, ZERO_CELL} \mbox{N * M,}
```

8.9.2.52 resize() [4/6]

```
el_rowsums resize (
            N ,
            ZERO_CELL )
```

8.9.2.53 resize() [5/6]

```
el resize ( \label{eq:n_* M_*, ZERO_CELL} \ )
```

8.9.2.54 resize() [6/6]

8.9.2.55 rm_cell() [1/3]

8.9.2.56 rm_cell() [2/3]

8.9.2.57 rm_cell() [3/3]

8.9.2.58 va_end()

```
va_end (
          args )
```

8.9.2.59 va_start()

```
va_start (
          args ,
          fmt )
```

8.9.2.60 vprintf()

8.9.3 Variable Documentation

8.9.3.1 add

```
uint const std::vector< uint > const std::vector< uint > bool add

Initial value:
{
    if (source.size() != target.size())
        throw std::length_error("-source- and -target- don't match on length.")
```

Definition at line 47 of file barraydense-meat.hpp.

8.9.3.2 ans

```
return ans
```

Definition at line 398 of file barraydense-meat.hpp.

8.9.3.3 check_bounds

```
bool check_bounds

Initial value:
{
    if (check_bounds)
    {
        out_of_range(i0,0u);
        out_of_range(i1,0u);
    }

for (uint j = 0u; j < M; ++j)
        std::swap(el[POS(i0, j)], el[POS(i1, j)])</pre>
```

Definition at line 654 of file barraydense-meat.hpp.

8.9.3.4 check_exists

```
uint bool int check_exists
Initial value:
{
    if (check_bounds)
        out_of_range(i,j)
```

Definition at line 655 of file barraydense-meat.hpp.

8.9.3.5 col

col

Definition at line 843 of file barraydense-meat.hpp.

8.9.3.6 const

const

Initial value:

```
if (i >= N)
     throw std::range_error("The row is out of range.")
```

Definition at line 360 of file barraydense-meat.hpp.

8.9.3.7 copy_data

```
bool copy_data
```

Definition at line 156 of file barraydense-meat.hpp.

8.9.3.8 data

```
data = data_
```

Definition at line 334 of file barraydense-meat.hpp.

8.9.3.9 delete_data

```
delete_data = delete_data_
```

Definition at line 335 of file barraydense-meat.hpp.

8.9.3.10 delete_data_

```
bool delete_data_
```

Initial value:

```
if ((data != nullptr) && delete_data)
     delete data
```

Definition at line 328 of file barraydense-meat.hpp.

8.9.3.11 el

```
return el == ZERO_CELL
```

Definition at line 381 of file barraydense-meat.hpp.

8.9.3.12 el_colsums

```
el_colsums[j] = (v.value - old)
```

Definition at line 675 of file barraydense-meat.hpp.

8.9.3.13 el_rowsums

```
el_rowsums[i] = (v.value - old)
```

Definition at line 674 of file barraydense-meat.hpp.

8.9.3.14 else

Definition at line 670 of file barraydense-meat.hpp.

8.9.3.15 false

false

Definition at line 767 of file barraydense-meat.hpp.

8.9.3.16 i1

```
uint il
```

Definition at line 721 of file barraydense-meat.hpp.

8.9.3.17 j

j

Definition at line 373 of file barraydense-meat.hpp.

8.9.3.18 j0

```
uint j0
```

Definition at line 720 of file barraydense-meat.hpp.

8.9.3.19 j1

```
uint j1
```

Definition at line 721 of file barraydense-meat.hpp.

8.9.3.20 M

```
M = M_{\underline{}}
```

Definition at line 57 of file barraydense-meat.hpp.

8.9.3.21 M_

```
uint M_
```

Initial value:

```
std::vector< Cell_Type > el_tmp(el)
```

Definition at line 43 of file barraydense-meat.hpp.

8.9.3.22 N

```
N = N_
```

Definition at line 56 of file barraydense-meat.hpp.

8.9.3.23 report

```
uint uint uint bool int int* report

Initial value:
{
    if (check_bounds) {
        out_of_range(i0, j0);
        out_of_range(i1, j1);
    }

    if (report != nullptr)
        (*report) = EXISTS::BOTH
```

Definition at line 724 of file barraydense-meat.hpp.

8.9.3.24 return

return

Definition at line 94 of file barraydense-meat.hpp.

8.9.3.25 source

```
uint const std::vector< uint >& source
```

Definition at line 44 of file barraydense-meat.hpp.

8.9.3.26 target

```
uint const std::vector< uint > const std::vector< uint >& target
```

Definition at line 45 of file barraydense-meat.hpp.

8.9.3.27 v

```
uint Cell_Type v
```

Definition at line 653 of file barraydense-meat.hpp.

8.9.3.28 val0

```
Cell_Type val0 = el[POS(i0,j0)]
```

Definition at line 742 of file barraydense-meat.hpp.

8.9.3.29 val1

```
Cell_Type val1 = el[POS(i1,j1)]
```

Definition at line 743 of file barraydense-meat.hpp.

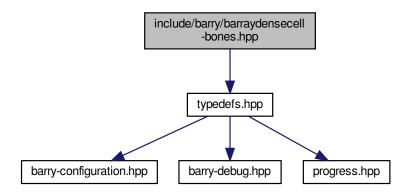
8.9.3.30 value

```
uint const std::vector< uint > const std::vector< uint > const std::vector< Cell_Type > & value
```

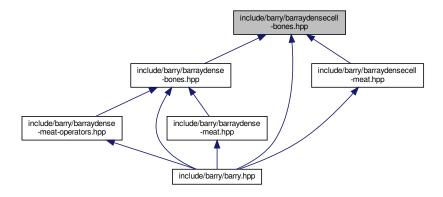
Definition at line 46 of file barraydense-meat.hpp.

8.10 include/barry/barraydensecell-bones.hpp File Reference

#include "typedefs.hpp"
Include dependency graph for barraydensecell-bones.hpp:



This graph shows which files directly or indirectly include this file:



Classes

class BArrayDenseCell
 Cell_Type, Data_Type

Macros

• #define POS(a, b) (a) + (b) * N

8.10.1 Macro Definition Documentation

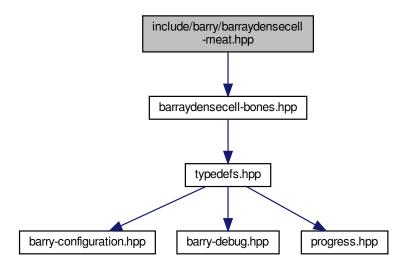
8.10.1.1 POS

```
#define POS(  a, \\ b \ ) \ (a) \ + \ (b) \ * \ N
```

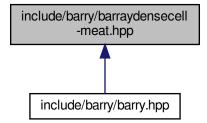
Definition at line 6 of file barraydensecell-bones.hpp.

8.11 include/barry/barraydensecell-meat.hpp File Reference

#include "barraydensecell-bones.hpp"
Include dependency graph for barraydensecell-meat.hpp:



This graph shows which files directly or indirectly include this file:



Macros

```
• #define POS(a, b) (a) + (b) * dat->N
```

8.11.1 Macro Definition Documentation

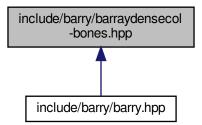
8.11.1.1 POS

```
#define POS(  a, \\ b ) \ (a) \ + \ (b) \ * \ dat -> N
```

Definition at line 6 of file barraydensecell-meat.hpp.

8.12 include/barry/barraydensecol-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



Classes

- class BArrayDenseCol< Cell_Type, Data_Type >
- class BArrayDenseCol_const< Cell_Type, Data_Type >

Macros

- #define POS(a, b) (b)*N + (a)
- #define POS_N(a, b, c) (b)*(c) + (a)
- #define ZERO_CELL static_cast<Cell_Type>(0.0)

8.12.1 Macro Definition Documentation

8.12.1.1 POS

```
#define POS(  a, \\ b ) (b)*N + (a)
```

Definition at line 4 of file barraydensecol-bones.hpp.

8.12.1.2 POS_N

Definition at line 5 of file barraydensecol-bones.hpp.

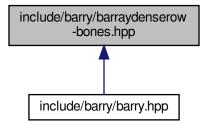
8.12.1.3 ZERO_CELL

```
#define ZERO_CELL static_cast<Cell_Type>(0.0)
```

Definition at line 6 of file barraydensecol-bones.hpp.

8.13 include/barry/barraydenserow-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



Classes

- class BArrayDenseRow< Cell_Type, Data_Type >
- class BArrayDenseRow_const< Cell_Type, Data_Type >

Macros

```
#define POS(a, b) (b) * N + (a)
#define POS_N(a, b, c) (b)*(c) + (a)
#define ZERO_CELL static_cast< Cell_Type >(0.0)
```

8.13.1 Macro Definition Documentation

8.13.1.1 POS

```
#define POS(  a, \\ b ) \ (b) \ * \ N \ + \ (a)
```

Definition at line 4 of file barraydenserow-bones.hpp.

8.13.1.2 POS N

Definition at line 5 of file barraydenserow-bones.hpp.

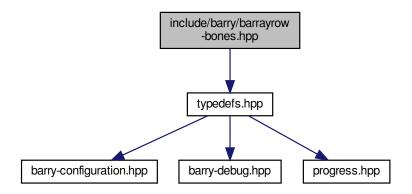
8.13.1.3 ZERO_CELL

```
#define ZERO_CELL static_cast< Cell_Type >(0.0)
```

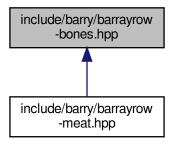
Definition at line 6 of file barraydenserow-bones.hpp.

8.14 include/barry/barrayrow-bones.hpp File Reference

#include "typedefs.hpp"
Include dependency graph for barrayrow-bones.hpp:



This graph shows which files directly or indirectly include this file:



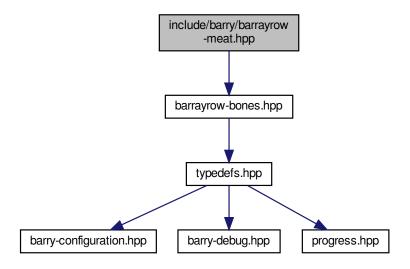
Classes

- class BArrayRow< Cell_Type, Data_Type >
- class BArrayRow_const< Cell_Type, Data_Type >

8.15 include/barry/barrayrow-meat.hpp File Reference

#include "barrayrow-bones.hpp"

Include dependency graph for barrayrow-meat.hpp:



Macros

- #define BARRY_BARRAYROW_MEAT_HPP 1
- #define BROW_TYPE() BArrayRow<Cell_Type, Data_Type>
- #define BROW_TEMPLATE_ARGS() < typename Cell_Type, typename Data_Type>
- #define BROW_TEMPLATE(a, b) template BROW_TEMPLATE_ARGS() inline a BROW_TYPE()::b

Functions

- BROW_TEMPLATE (void, operator=)(const BROW_TYPE() &val)
- BROW_TEMPLATE (void, operator+=)(const BROW_TYPE() &val)
- BROW_TEMPLATE (void, operator-=)(const BROW_TYPE() &val)
- BROW_TEMPLATE (void, operator*=)(const BROW_TYPE() &val)
- BROW TEMPLATE (void, operator/=)(const BROW TYPE() &val)

8.15.1 Macro Definition Documentation

8.15.1.1 BARRY BARRAYROW MEAT HPP

#define BARRY_BARRAYROW_MEAT_HPP 1

Definition at line 4 of file barrayrow-meat.hpp.

8.15.1.2 BROW_TEMPLATE

Definition at line 10 of file barrayrow-meat.hpp.

8.15.1.3 BROW_TEMPLATE_ARGS

```
#define BROW_TEMPLATE_ARGS() <typename Cell_Type, typename Data_Type>
```

Definition at line 8 of file barrayrow-meat.hpp.

8.15.1.4 BROW_TYPE

```
#define BROW_TYPE( ) BArrayRow<Cell_Type, Data_Type>
```

Definition at line 6 of file barrayrow-meat.hpp.

8.15.2 Function Documentation

8.15.2.1 BROW_TEMPLATE() [1/5]

Definition at line 47 of file barrayrow-meat.hpp.

8.15.2.2 BROW_TEMPLATE() [2/5]

Definition at line 27 of file barrayrow-meat.hpp.

8.15.2.3 BROW_TEMPLATE() [3/5]

Definition at line 36 of file barrayrow-meat.hpp.

8.15.2.4 BROW_TEMPLATE() [4/5]

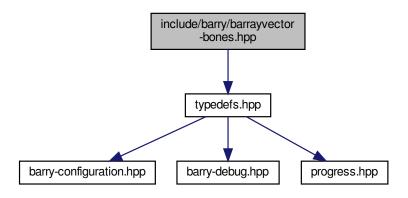
Definition at line 57 of file barrayrow-meat.hpp.

8.15.2.5 BROW_TEMPLATE() [5/5]

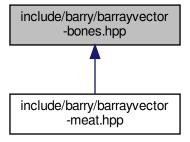
Definition at line 13 of file barrayrow-meat.hpp.

8.16 include/barry/barrayvector-bones.hpp File Reference

```
#include "typedefs.hpp"
Include dependency graph for barrayvector-bones.hpp:
```



This graph shows which files directly or indirectly include this file:

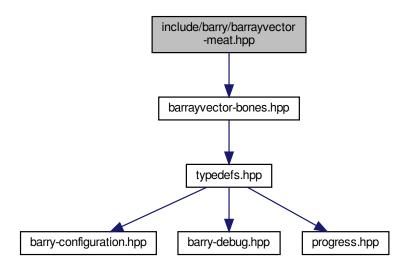


Classes

- class BArrayVector< Cell_Type, Data_Type >
 Row or column of a BArray
- class BArrayVector_const< Cell_Type, Data_Type >

8.17 include/barry/barrayvector-meat.hpp File Reference

#include "barrayvector-bones.hpp"
Include dependency graph for barrayvector-meat.hpp:



Macros

• #define BARRY BARRAYVECTOR MEAT HPP 1

8.17.1 Macro Definition Documentation

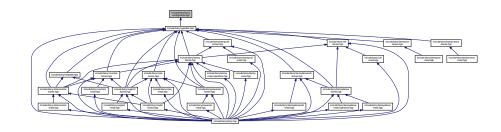
8.17.1.1 BARRY_BARRAYVECTOR_MEAT_HPP

```
#define BARRY_BARRAYVECTOR_MEAT_HPP 1
```

Definition at line 4 of file barrayvector-meat.hpp.

8.18 include/barry/barry-configuration.hpp File Reference

This graph shows which files directly or indirectly include this file:



Configuration MACROS

These are mostly related to performance. The definitions follow:

- BARRY_USE_UNORDERED_MAP If specified, then barry is compiled using std::unordered_map. Otherwise it will use std::map for the arrays.
- BARRY_USE_SAFE_EXP When specified, it will multiply all likelihoods in Model by (1/-100)/(1/-100) so that numerical overflows are avoided.
- BARRY_USE_ISFINITE When specified, it will introduce a macro that checks whether the likelihood is finite or not.
- printf_barry If not specified, will be defined as printf.
- BARRY_DEBUG_LEVEL, when defined, will make things verbose.
- #define BARRY_SAFE_EXP -100.0
- #define BARRY_ISFINITE(a)
- #define BARRY CHECK SUPPORT(x, maxs)
- #define printf barry printf
- #define BARRY_MAX_NUM_ELEMENTS static_cast< size_t >(UINT_MAX/2u)
- template<typename Ta , typename Tb >
 using Map = std::map< Ta, Tb >

8.18.1 Macro Definition Documentation

8.18.1.1 BARRY_CHECK_SUPPORT

```
#define BARRY_CHECK_SUPPORT(
          x,
          maxs )
```

Definition at line 47 of file barry-configuration.hpp.

8.18.1.2 BARRY_ISFINITE

Definition at line 40 of file barry-configuration.hpp.

8.18.1.3 BARRY_MAX_NUM_ELEMENTS

```
#define BARRY_MAX_NUM_ELEMENTS static_cast< size_t >(UINT_MAX/2u)
```

Definition at line 55 of file barry-configuration.hpp.

8.18.1.4 BARRY_SAFE_EXP

```
#define BARRY_SAFE_EXP -100.0
```

Definition at line 33 of file barry-configuration.hpp.

8.18.1.5 printf_barry

```
#define printf_barry printf
```

Definition at line 51 of file barry-configuration.hpp.

8.18.2 Typedef Documentation

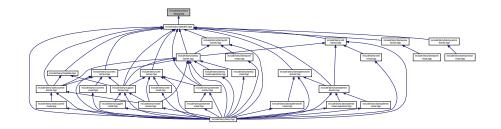
8.18.2.1 Map

```
template<typename Ta , typename Tb >
using Map = std::map<Ta,Tb>
```

Definition at line 27 of file barry-configuration.hpp.

8.19 include/barry/barry-debug.hpp File Reference

This graph shows which files directly or indirectly include this file:



Macros

• #define BARRY_DEBUG_LEVEL 0

8.19.1 Macro Definition Documentation

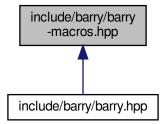
8.19.1.1 BARRY_DEBUG_LEVEL

#define BARRY_DEBUG_LEVEL 0

Definition at line 5 of file barry-debug.hpp.

8.20 include/barry/barry-macros.hpp File Reference

This graph shows which files directly or indirectly include this file:



Macros

- #define BARRY_ZERO Cell<Cell_Type>(0.0)
- #define BARRY_ZERO_DENSE static_cast<Cell_Type>(0.0)
- #define BARRY_ONE Cell<Cell_Type>(1.0)
- #define BARRY_ONE_DENSE static_cast<Cell_Type>(1.0)
- #define BARRY_UNUSED(expr) do { (void)(expr); } while (0);

8.20.1 Macro Definition Documentation

8.20.1.1 BARRY_ONE

```
#define BARRY_ONE CellCell_Type>(1.0)
```

Definition at line 7 of file barry-macros.hpp.

8.20.1.2 BARRY_ONE_DENSE

```
#define BARRY_ONE_DENSE static_cast<Cell_Type>(1.0)
```

Definition at line 8 of file barry-macros.hpp.

8.20.1.3 BARRY_UNUSED

Definition at line 10 of file barry-macros.hpp.

8.20.1.4 BARRY_ZERO

```
#define BARRY_ZERO Cell<Cell_Type>(0.0)
```

Definition at line 4 of file barry-macros.hpp.

8.20.1.5 BARRY_ZERO_DENSE

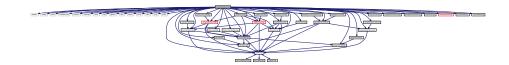
```
#define BARRY_ZERO_DENSE static_cast<Cell_Type>(0.0)
```

Definition at line 5 of file barry-macros.hpp.

8.21 include/barry/barry.hpp File Reference

```
#include <iostream>
#include <cstdarg>
#include <vector>
#include <unordered_map>
#include <functional>
#include <stdexcept>
#include <cmath>
#include <map>
#include <algorithm>
#include <utility>
#include <random>
#include <climits>
#include <cfloat>
#include <string>
#include <cstdint>
#include "typedefs.hpp"
#include "barry-macros.hpp"
#include "cell-bones.hpp"
#include "cell-meat.hpp"
#include "barray-bones.hpp"
#include "barraycell-bones.hpp"
#include "barray-meat.hpp"
#include "barraycell-meat.hpp"
#include "barray-meat-operators.hpp"
#include "barraydense-bones.hpp"
#include "barraydensecell-bones.hpp"
#include "barraydenserow-bones.hpp"
#include "barraydensecol-bones.hpp"
#include "barraydense-meat.hpp"
#include "barraydensecell-meat.hpp"
#include "barraydense-meat-operators.hpp"
#include "counters-bones.hpp"
#include "counters-meat.hpp"
#include "statscounter-bones.hpp"
#include "statscounter-meat.hpp"
#include "support-bones.hpp"
#include "support-meat.hpp"
#include "powerset-bones.hpp"
#include "powerset-meat.hpp"
#include "model-bones.hpp"
#include "model-meat.hpp"
#include "rules-bones.hpp"
#include "rules-meat.hpp"
#include "counters/network.hpp"
#include "counters/phylo.hpp"
```

#include "counters/defm.hpp"
Include dependency graph for barry.hpp:



Namespaces

barry

barry: Your go-to motif accountant

• barry::counters

Tree class and Treelterator class.

- barry::counters::network
- · barry::counters::phylo
- barry::counters::defm

Macros

- #define BARRY HPP
- #define BARRY_VERSION 0.1
- #define COUNTER_FUNCTION(a)
- #define COUNTER_LAMBDA(a)
- #define RULE_FUNCTION(a)
- #define RULE_LAMBDA(a)

8.21.1 Macro Definition Documentation

8.21.1.1 BARRY_HPP

#define BARRY_HPP

Definition at line 22 of file barry.hpp.

8.21.1.2 BARRY_VERSION

#define BARRY_VERSION 0.1

Definition at line 24 of file barry.hpp.

8.21.1.3 COUNTER_FUNCTION

Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
inline double (a) (const Array_Type & Array, uint i, uint j, Data_Type & data)
```

Definition at line 89 of file barry.hpp.

8.21.1.4 COUNTER_LAMBDA

Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
Counter_fun_type<Array_Type, Data_Type> a = \
[](const Array_Type & Array, uint i, uint j, Data_Type & data)
```

Definition at line 92 of file barry.hpp.

8.21.1.5 RULE FUNCTION

Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
inline bool (a) (const Array_Type & Array, uint i, uint j, Data_Type & data) \
```

Definition at line 96 of file barry.hpp.

8.21.1.6 RULE LAMBDA

```
#define RULE_LAMBDA( a )
```

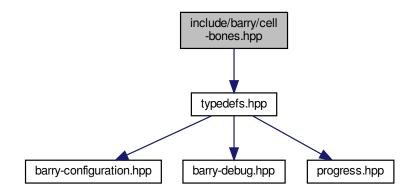
Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
Rule_fun_type<Array_Type, Data_Type> a = \
[](const Array_Type & Array, uint i, uint j, Data_Type & data)
```

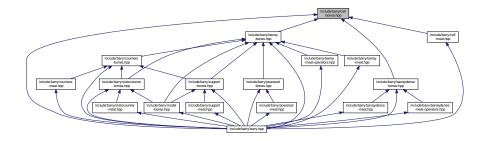
Definition at line 99 of file barry.hpp.

8.22 include/barry/cell-bones.hpp File Reference

#include "typedefs.hpp"
Include dependency graph for cell-bones.hpp:



This graph shows which files directly or indirectly include this file:



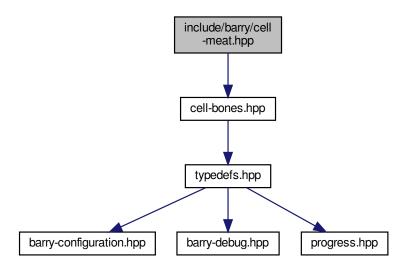
Classes

class Cell
 Cell_Type >
 Entries in BArray. For now, it only has two members:

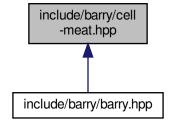
8.23 include/barry/cell-meat.hpp File Reference

#include "cell-bones.hpp"

Include dependency graph for cell-meat.hpp:



This graph shows which files directly or indirectly include this file:



8.24 include/barry/col-bones.hpp File Reference

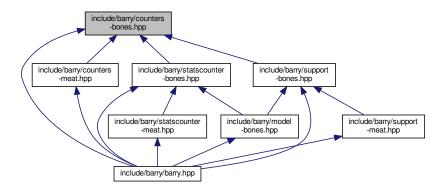
8.25 include/barry/counters-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "barray-bones.hpp"
```

Include dependency graph for counters-bones.hpp:



This graph shows which files directly or indirectly include this file:

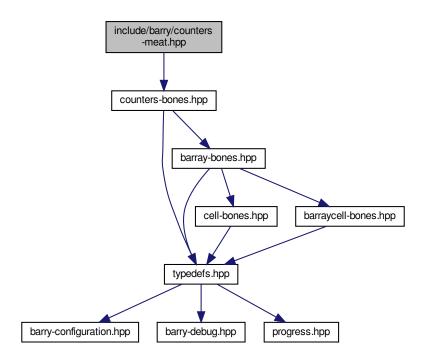


Classes

- class Counters
 Array_Type, Data_Type >
 Vector of counters.

8.26 include/barry/counters-meat.hpp File Reference

#include "counters-bones.hpp"
Include dependency graph for counters-meat.hpp:



This graph shows which files directly or indirectly include this file:



Macros

- #define COUNTER_TYPE() Counter<Array_Type,Data_Type>
- #define COUNTER_TEMPLATE_ARGS() < typename Array_Type, typename Data_Type>

- #define COUNTER_TEMPLATE(a, b) template COUNTER_TEMPLATE_ARGS() inline a COUNTER_TYPE() ↔ ::b
- #define COUNTERS_TYPE() Counters<Array_Type,Data_Type>
- #define COUNTERS_TEMPLATE_ARGS() < typename Array_Type, typename Data_Type>

Functions

- COUNTER_TEMPLATE (, Counter)(const Counter< Array_Type
- Data Type init fun (counter .init fun)
- Data Type &&counter init fun (std::move(counter .init fun))
- Data Type &&counter data (std::move(counter .data))
- Data_Type &&counter_ name (std::move(counter_.name))
- Data_Type &&counter_ desc (std::move(counter_.desc))

Move constructor.

- COUNTER TEMPLATE (COUNTER TYPE(), operator=)(const Counter< Array Type
- COUNTER_TEMPLATE (COUNTER_TYPE() &, operator=)(Counter< Array_Type
- COUNTER TEMPLATE (double, count)(Array Type & Array

< Move assignment

- return count_fun (Array, i, j, data)
- COUNTER_TEMPLATE (double, init)(Array_Type &Array
- return init_fun (Array, i, j, data)
- COUNTER_TEMPLATE (std::string, get_name)() const
- COUNTER_TEMPLATE (std::string, get_description)() const
- COUNTERS_TEMPLATE (, Counters)()
- COUNTERS_TEMPLATE (COUNTER_TYPE() &, operator[])(uint idx)
- Data_Type Counters (Counters < Array_Type, Data_Type > &&counters_) noexcept
- COUNTERS_TEMPLATE (COUNTERS_TYPE(), operator=)(const Counters < Array_Type
- COUNTERS_TEMPLATE (COUNTERS_TYPE() &, operator=)(Counters< Array_Type
- COUNTERS_TEMPLATE (void, add_counter)(Counter< Array_Type
- COUNTERS_TEMPLATE (std::vector< std::string >, get_names)() const
- COUNTERS_TEMPLATE (std::vector< std::string >, get_descriptions)() const

Variables

- Data Type & counter
- Data_Type &&counter_ noexcept
- uint i
- · uint uint j
- return * this
- Data_Type counter
- return
- Data_Type count_fun_
- Data_Type Counter_fun_type
 Array_Type, Data_Type > init_fun_
- Data_Type Counter_fun_type
 Array_Type, Data_Type > Data_Type data_
- Data_Type Counter_fun_type
 Array_Type, Data_Type > Data_Type std::string name_
- Data_Type Counter_fun_type
 Array_Type, Data_Type > Data_Type std::string std::string desc_

8.26.1 Macro Definition Documentation

8.26.1.1 COUNTER_TEMPLATE

Definition at line 10 of file counters-meat.hpp.

8.26.1.2 COUNTER_TEMPLATE_ARGS

```
#define COUNTER_TEMPLATE_ARGS() <typename Array_Type, typename Data_Type>
```

Definition at line 8 of file counters-meat.hpp.

8.26.1.3 COUNTER_TYPE

```
#define COUNTER_TYPE( ) Counter<Array_Type, Data_Type>
```

Definition at line 6 of file counters-meat.hpp.

8.26.1.4 COUNTERS_TEMPLATE

Definition at line 118 of file counters-meat.hpp.

8.26.1.5 COUNTERS_TEMPLATE_ARGS

```
#define COUNTERS_TEMPLATE_ARGS() <typename Array_Type, typename Data_Type>
```

Definition at line 116 of file counters-meat.hpp.

8.26.1.6 COUNTERS_TYPE

```
#define COUNTERS_TYPE( ) Counters<Array_Type,Data_Type>
```

Definition at line 114 of file counters-meat.hpp.

8.26.2 Function Documentation

8.26.2.1 count fun()

8.26.2.2 COUNTER_TEMPLATE() [1/7]

```
COUNTER_TEMPLATE (

Counter ) const
```

8.26.2.3 COUNTER_TEMPLATE() [2/7]

8.26.2.4 COUNTER_TEMPLATE() [3/7]

8.26.2.5 COUNTER_TEMPLATE() [4/7]

< Move assignment

8.26.2.6 COUNTER_TEMPLATE() [5/7]

```
COUNTER_TEMPLATE ( \label{eq:counter} \mbox{double ,} \\ \mbox{init ) } \&
```

8.26.2.7 COUNTER_TEMPLATE() [6/7]

```
COUNTER_TEMPLATE (
          std::string ,
          get_description ) const
```

Definition at line 106 of file counters-meat.hpp.

8.26.2.8 COUNTER_TEMPLATE() [7/7]

```
COUNTER_TEMPLATE (
          std::string ,
          get_name ) const
```

Definition at line 102 of file counters-meat.hpp.

8.26.2.9 Counters()

Definition at line 132 of file counters-meat.hpp.

8.26.2.10 COUNTERS_TEMPLATE() [1/7]

```
COUNTERS_TEMPLATE (

Counters )
```

Definition at line 121 of file counters-meat.hpp.

8.26.2.11 COUNTERS_TEMPLATE() [2/7]

Definition at line 123 of file counters-meat.hpp.

8.26.2.12 COUNTERS_TEMPLATE() [3/7]

```
COUNTERS_TEMPLATE (

COUNTERS_TYPE() & ,

operator )
```

8.26.2.13 COUNTERS_TEMPLATE() [4/7]

```
COUNTERS_TEMPLATE (

COUNTERS_TYPE() ,

operator ) const
```

8.26.2.14 COUNTERS_TEMPLATE() [5/7]

```
COUNTERS_TEMPLATE (
          std::vector< std::string > ,
          get_descriptions ) const
```

Definition at line 194 of file counters-meat.hpp.

8.26.2.15 COUNTERS_TEMPLATE() [6/7]

```
COUNTERS_TEMPLATE (
          std::vector< std::string > ,
          get_names ) const
```

Definition at line 183 of file counters-meat.hpp.

8.26.2.16 COUNTERS_TEMPLATE() [7/7]

```
COUNTERS_TEMPLATE (
     void ,
     add_counter )
```

8.26.2.17 data()

8.26.2.18 desc()

Move constructor.

Definition at line 33 of file counters-meat.hpp.

8.26.2.19 init_fun() [1/3]

8.26.2.20 init_fun() [2/3]

Definition at line 15 of file counters-meat.hpp.

8.26.2.21 init_fun() [3/3]

8.26.2.22 name()

8.26.3 Variable Documentation

8.26.3.1 count_fun_

```
Data_Type count_fun_
```

Definition at line 163 of file counters-meat.hpp.

8.26.3.2 counter

```
Data_Type counter

Initial value:
{
    data.push_back(counter)
```

Definition at line 154 of file counters-meat.hpp.

8.26.3.3 counter_

```
Data_Type & counter_

Initial value:
{
    if (this != &counter_) {
        this->count_fun = counter_.count_fun;
        this->init_fun = counter_.init_fun;

        this->data = counter_.data;
        this->name = counter_.name;
        this->desc = counter_.desc;
    }
    return *this
```

Definition at line 14 of file counters-meat.hpp.

8.26.3.4 data_

```
Data_Type Counter_fun_type<Array_Type,Data_Type> Data_Type data_
```

Definition at line 165 of file counters-meat.hpp.

8.26.3.5 desc

Data_Type Counter_fun_type<Array_Type,Data_Type > Data_Type std::string std::string desc_

Initial value:

```
{
```

```
data.push_back(Counter<Array_Type,Data_Type>(
    count_fun_,
    init_fun_,
    data_,
    name_,
    desc_
```

Definition at line 167 of file counters-meat.hpp.

8.26.3.6 i

uint i

Definition at line 82 of file counters-meat.hpp.

8.26.3.7 init_fun_

```
Data_Type Counter_fun_type<Array_Type,Data_Type> init_fun_
```

Definition at line 164 of file counters-meat.hpp.

8.26.3.8 j

```
uint uint j
```

Initial value:

```
if (count_fun == nullptr)
    return 0.0
```

Definition at line 82 of file counters-meat.hpp.

8.26.3.9 name_

```
Data_Type Counter_fun_type<Array_Type,Data_Type> Data_Type std::string name_
Definition at line 166 of file counters-meat.hpp.
```

8.26.3.10 noexcept

```
Data_Type &&counters_ noexcept

Initial value:
{
    if (this != &counter_)
    {
        this->data = std::move(counter_.data);

        this->init_fun = std::move(counter_.count_fun);
        this->name = std::move(counter_.init_fun);

        this->desc = std::move(counter_.name);
        this->desc = std::move(counter_.desc);
    }
    return *this
```

Definition at line 28 of file counters-meat.hpp.

8.26.3.11 return

return

Definition at line 159 of file counters-meat.hpp.

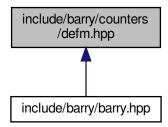
8.26.3.12 this

```
return* this
```

Definition at line 140 of file counters-meat.hpp.

8.27 include/barry/counters/defm.hpp File Reference

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 uint or double vectors.

Macros

Macros for defining counters

- #define DEFM COUNTER(a)
- #define DEFM_COUNTER_LAMBDA(a)

Macros for defining rules

- #define DEFM RULE(a)
- #define DEFM_RULE_LAMBDA(a)

Typedefs

Convenient typedefs for network objects.

```
    typedef BArrayDense < int, DEFMData > DEFMArray
    template < typename Tarray = DEFMArray > using DEFMCounter = Counter < Tarray, DEFMCounterData >
    template < typename Tarray = DEFMArray > using DEFMCounters = Counters < Tarray, DEFMCounterData >
    template < typename Tarray = DEFMArray > using DEFMSupport = Support < Tarray, DEFMCounterData >
    template < typename Tarray = DEFMArray > using DEFMStatsCounter = StatsCounter < Tarray, DEFMCounterData >
    template < typename Tarray > using DEFMModel = Model < Tarray, DEFMCounterData >
    template < typename Tarray = DEFMArray > using DEFMRule = Rule < Tarray, bool >
    template < typename Tarray = DEFMArray > using DEFMRules = Rules < Tarray, bool >
```

Functions

Rules for network models

Parameters

```
rules | A pointer to a DEFMRules object (Rules < DEFMArray, bool > ).
```

```
    template < typename Tarray = DEFMArray >
    void rules_zerodiag (DEFMRules < Tarray > *rules)
    Number of edges.
```

8.27.1 Macro Definition Documentation

8.27.1.1 DEFM_COUNTER

Function for definition of a network counter function

Definition at line 125 of file defm.hpp.

8.27.1.2 DEFM_COUNTER_LAMBDA

Lambda function for definition of a network counter function

Definition at line 130 of file defm.hpp.

8.27.1.3 DEFM_RULE

Function for definition of a network counter function

Definition at line 141 of file defm.hpp.

8.27.1.4 DEFM_RULE_LAMBDA

```
#define DEFM_RULE_LAMBDA( \it a )
```

Value:

```
Rule_fun_type<Tarray, bool> a = \
[](const Tarray & Array, uint i, uint j, bool & data)
```

Lambda function for definition of a network counter function

Definition at line 146 of file defm.hpp.

8.27.2 Typedef Documentation

8.27.2.1 **DEFMArray**

```
typedef BArrayDense<int, DEFMData> DEFMArray
```

Definition at line 97 of file defm.hpp.

8.27.2.2 DEFMCounter

```
template<typename Tarray = DEFMArray>
using DEFMCounter = Counter<Tarray, DEFMCounterData >
```

Definition at line 100 of file defm.hpp.

8.27.2.3 DEFMCounters

```
template<typename Tarray = DEFMArray>
using DEFMCounters = Counters<Tarray, DEFMCounterData>
```

Definition at line 103 of file defm.hpp.

8.27.2.4 DEFMModel

```
template<typename Tarray >
using DEFMModel = Model<Tarray, DEFMCounterData>
```

Definition at line 112 of file defm.hpp.

8.27.2.5 **DEFMRule**

```
template<typename Tarray = DEFMArray>
using DEFMRule = Rule<Tarray, bool>
```

Definition at line 115 of file defm.hpp.

8.27.2.6 **DEFMRules**

```
template<typename Tarray = DEFMArray>
using DEFMRules = Rules<Tarray, bool>
```

Definition at line 118 of file defm.hpp.

8.27.2.7 DEFMStatsCounter

```
template<typename Tarray = DEFMArray>
using DEFMStatsCounter = StatsCounter<Tarray, DEFMCounterData>
```

Definition at line 109 of file defm.hpp.

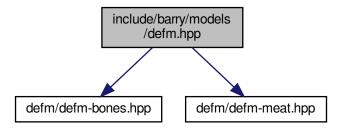
8.27.2.8 DEFMSupport

```
template<typename Tarray = DEFMArray>
using DEFMSupport = Support<Tarray, DEFMCounterData >
```

Definition at line 106 of file defm.hpp.

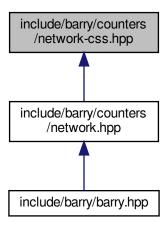
8.28 include/barry/models/defm.hpp File Reference

```
#include "defm/defm-bones.hpp"
#include "defm/defm-meat.hpp"
Include dependency graph for defm.hpp:
```



8.29 include/barry/counters/network-css.hpp File Reference

This graph shows which files directly or indirectly include this file:



Macros

- #define CSS_SIZE()
- #define CSS_CASE_TRUTH() if ((i < n) && (j < n))
- #define CSS_TRUE_CELLS()
- #define CSS_CASE_PERCEIVED() else if ((($i \ge s$) && (i < e)) & (($j \ge s$) && (j < e)))
- #define CSS PERCEIVED CELLS()
- #define CSS CASE ELSE()
- #define CSS_CHECK_SIZE_INIT()
- #define CSS_CHECK_SIZE()
- #define CSS_APPEND(name)
- #define CSS_NET_COUNTER_LAMBDA_INIT()

Functions

• template<typename Tnet = Network> void counter_css_partially_false_recip_commi (NetCounters< Tnet > *counters, uint netsize, const std ← ::vector< uint > &end_)

Counts errors of commission.

• template<typename Tnet = Network> void counter_css_partially_false_recip_omiss (NetCounters< Tnet > *counters, uint netsize, const std
::vector< uint > &end)

Counts errors of omission.

• template<typename Tnet = Network> void counter_css_completely_false_recip_comiss (NetCounters< Tnet > *counters, uint netsize, const std ← ::vector< uint > &end_)

Counts completely false reciprocity (comission)

```
• template<typename Tnet = Network>
  void counter_css_completely_false_recip_omiss (NetCounters< Tnet > *counters, uint netsize, const std↔
  ::vector< uint > &end_)
     Counts completely false reciprocity (omission)
• template<typename Tnet = Network>
  void counter css mixed recip (NetCounters< Tnet > *counters, uint netsize, const std::vector< uint >
  &end )
     Counts mixed reciprocity errors.
• template<typename Tnet = Network>
 void counter_css_census01 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
• template<typename Tnet = Network>
 void counter css census02 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
template<typename Tnet = Network>
 void counter_css_census03 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
template<typename Tnet = Network>
  void counter_css_census04 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
 _)
• template<typename Tnet = Network>
 void counter css census05 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
template<typename Tnet = Network>
  void counter css census06 (NetCounters< Tnet > *counters, uint netsize, const std::vector< uint > &end↔
• template<typename Tnet = Network>
 void counter_css_census07 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
 _)
• template<typename Tnet = Network>
  void counter_css_census08 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
 _)
• template<typename Tnet = Network>
 void counter css census09 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←

    template<typename Tnet = Network>

  void counter_css_census10 (NetCounters< Tnet > *counters, uint netsize, const std::vector< uint > &end↔
  _)
```

8.29.1 Macro Definition Documentation

8.29.1.1 CSS_APPEND

Definition at line 42 of file network-css.hpp.

8.29.1.2 CSS_CASE_ELSE

```
#define CSS_CASE_ELSE( )
```

Definition at line 27 of file network-css.hpp.

8.29.1.3 CSS_CASE_PERCEIVED

```
      \# define \ CSS\_CASE\_PERCEIVED ( ) \ else \ if \ (((i >= s) \ \&\& \ (i < e)) \ \& \ ((j >= s) \ \&\& \ (j < e)))
```

Definition at line 20 of file network-css.hpp.

8.29.1.4 CSS CASE TRUTH

```
#define CSS_CASE_TRUTH( ) if ((i < n) && (j < n))
```

Definition at line 13 of file network-css.hpp.

8.29.1.5 CSS_CHECK_SIZE

```
#define CSS_CHECK_SIZE( )
```

Value:

```
for (uint i = 0u; i < end_.size(); ++i) {\
   if (i == 0u) continue; \
   else if (end_[i] < end_[i-lu]) \
       throw std::logic_error("Endpoints should be specified in order.");}</pre>
```

Definition at line 37 of file network-css.hpp.

8.29.1.6 CSS_CHECK_SIZE_INIT

```
#define CSS_CHECK_SIZE_INIT( )
```

Value:

```
/* The indices fall within the network */ \
if ((data.indices.at(0) > Array.ncol()) \
| (data.indices.at(2) > Array.ncol())) \
throw std::range_error("The network does not match the prescribed size.");
```

Definition at line 31 of file network-css.hpp.

8.29.1.7 CSS_NET_COUNTER_LAMBDA_INIT

Definition at line 49 of file network-css.hpp.

8.29.1.8 CSS_PERCEIVED_CELLS

```
#define CSS_PERCEIVED_CELLS()

Value:
    double tji = static_cast<double>(Array(j - s, i - s, false)); \
    double pji = static_cast<double>(Array(j, i, false)); \
    double tij = static_cast<double>(Array(i - s, j - s, false));
```

Definition at line 21 of file network-css.hpp.

8.29.1.9 CSS_SIZE

```
#define CSS_SIZE()

Value:
    uint n = data.indices[0u]; \
    uint s = data.indices[1u]; \
    uint e = data.indices[2u];
```

Definition at line 7 of file network-css.hpp.

8.29.1.10 CSS_TRUE_CELLS

```
#define CSS_TRUE_CELLS( )

Value:
    double tji = static_cast<double>(Array(j, i, false)); \
    double pij = static_cast<double>(Array(i + s, j + s, false)); \
    double pji = static_cast<double>(Array(j + s, i + s, false));
```

Definition at line 14 of file network-css.hpp.

8.29.2 Function Documentation

8.29.2.1 counter_css_census01()

Definition at line 275 of file network-css.hpp.

8.29.2.2 counter_css_census02()

Definition at line 325 of file network-css.hpp.

8.29.2.3 counter_css_census03()

Definition at line 364 of file network-css.hpp.

8.29.2.4 counter_css_census04()

Definition at line 403 of file network-css.hpp.

8.29.2.5 counter_css_census05()

Definition at line 442 of file network-css.hpp.

8.29.2.6 counter_css_census06()

Definition at line 481 of file network-css.hpp.

8.29.2.7 counter_css_census07()

Definition at line 520 of file network-css.hpp.

8.29.2.8 counter_css_census08()

Definition at line 559 of file network-css.hpp.

8.29.2.9 counter_css_census09()

Definition at line 598 of file network-css.hpp.

8.29.2.10 counter_css_census10()

Definition at line 637 of file network-css.hpp.

8.29.2.11 counter_css_completely_false_recip_comiss()

Counts completely false reciprocity (comission)

Definition at line 154 of file network-css.hpp.

8.29.2.12 counter_css_completely_false_recip_omiss()

Counts completely false reciprocity (omission)

Definition at line 194 of file network-css.hpp.

8.29.2.13 counter_css_mixed_recip()

Counts mixed reciprocity errors.

Definition at line 234 of file network-css.hpp.

8.29.2.14 counter_css_partially_false_recip_commi()

Counts errors of commission.

Parameters

netsize	Size of the reference (true) network
end←	Vector indicating one past the ending index of each network. (see details)
_	

The end_ parameter should be of length ${\tt N}$ of networks - 1. It is assumed that the first network ends at netsize.

Definition at line 63 of file network-css.hpp.

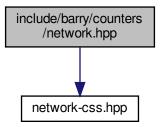
8.29.2.15 counter_css_partially_false_recip_omiss()

Counts errors of omission.

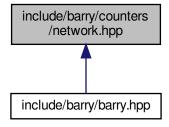
Definition at line 110 of file network-css.hpp.

8.30 include/barry/counters/network.hpp File Reference

#include "network-css.hpp"
Include dependency graph for network.hpp:



This graph shows which files directly or indirectly include this file:



Classes

class NetworkData

Data class for Networks.

· class NetCounterData

Data class used to store arbitrary uint or double vectors.

Macros

- #define NET_C_DATA_IDX(i) (data.indices[i])
- #define NET_C_DATA_NUM(i) (data.numbers[i])

Macros for defining counters

- #define NETWORK COUNTER(a)
- #define NETWORK COUNTER LAMBDA(a)
- #define NETWORKDENSE_COUNTER_LAMBDA(a)

Macros for defining rules

- #define NETWORK RULE(a)
- #define NETWORK_RULE_LAMBDA(a)

Functions

```
template<typename Tnet = Network>
  void counter_edges (NetCounters < Tnet > *counters)
     Number of edges.
• template<typename Tnet = Network>
  void counter isolates (NetCounters < Tnet > *counters)
     Number of isolated vertices.

    template<> void counter_isolates (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter mutual (NetCounters< Tnet > *counters)
     Number of mutual ties.
• template<typename Tnet = Network>
  void counter_istar2 (NetCounters < Tnet > *counters)

    template<> void counter_istar2 (NetCounters< NetworkDense > *counters)

    template<typename Tnet = Network>

  void counter ostar2 (NetCounters< Tnet > *counters)

    template<> void counter ostar2 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter_ttriads (NetCounters < Tnet > *counters)

    template<> void counter_ttriads (NetCounters< NetworkDense > *counters)

template<typename Tnet = Network>
  void counter ctriads (NetCounters< Tnet > *counters)

    template<> void counter ctriads (NetCounters< NetworkDense > *counters)

template<typename Tnet = Network>
  void counter_density (NetCounters < Tnet > *counters)
• template<typename Tnet = Network>
  void counter_idegree15 (NetCounters< Tnet > *counters)

    template<> void counter idegree15 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter odegree15 (NetCounters< Tnet > *counters)

    template<> void counter_odegree15 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter_absdiff (NetCounters< Tnet > *counters, uint attr_id, double alpha=1.0)
     Sum of absolute attribute difference between ego and alter.
• template<typename Tnet = Network>
  void counter_diff (NetCounters < Tnet > *counters, uint attr_id, double alpha=1.0, double tail_head=true)
     Sum of attribute difference between ego and alter to pow(alpha)

    NETWORK COUNTER (init single attr)
```

• template<typename Tnet = Network>

• template<typename Tnet = Network>

void counter_nodeicov (NetCounters< Tnet > *counters, uint attr_id)

void counter_nodeocov (NetCounters< Tnet > *counters, uint attr_id)

```
    template<typename Tnet = Network>
        void counter_nodecov (NetCounters< Tnet > *counters, uint attr_id)
    template<typename Tnet = Network>
        void counter_nodematch (NetCounters< Tnet > *counters, uint attr_id)
    template<typename Tnet = Network>
        void counter_idegree (NetCounters< Tnet > *counters, std::vector< uint > d)
        Counts number of vertices with a given in-degree.
    template<>> void counter_idegree (NetCounters< NetworkDense > *counters, std::vector< uint > d)
    template<typename Tnet = Network>
        void counter_odegree (NetCounters< Tnet > *counters, std::vector< uint > d)
    Counts number of vertices with a given out-degree.
    template<>> void counter_odegree (NetCounters< NetworkDense > *counters, std::vector< uint > d)
    template<typename Tnet = Network>
        void counter_degree (NetCounters< Tnet > *counters, std::vector< uint > d)
    template<typename Tnet = Network>
        void counter_degree (NetCounters< Tnet > *counters, std::vector< uint > d)
    Counts number of vertices with a given out-degree.
```

Rules for network models

Parameters

```
rules | A pointer to a NetRules object (Rules < Network, bool > ).
```

template < typename Tnet = Network >
 void rules_zerodiag (NetRules < Tnet > *rules)
 Number of edges.

Convenient typedefs for network objects.

- #define BARRY_ZERO_NETWORK 0.0
- #define BARRY_ZERO_NETWORK_DENSE 0
- typedef BArray< double, NetworkData > Network
- typedef BArrayDense< int, NetworkData > NetworkDense
- template<typename Tnet = Network>
 using NetCounter = Counter< Tnet, NetCounterData >
- template < typename Tnet = Network >
 using NetCounters = Counters < Tnet, NetCounterData >
- template<typename Tnet = Network>
 using NetSupport = Support< Tnet, NetCounterData >
- template < typename Tnet = Network >
 using NetStatsCounter = StatsCounter < Tnet, NetCounterData >
- template < typename Tnet >
 using NetModel = Model < Tnet, NetCounterData >
- template<typename Tnet = Network>
 using NetRule = Rule
 Tnet, bool >
- template<typename Tnet = Network>
 using NetRules = Rules< Tnet, bool >

8.30.1 Macro Definition Documentation

8.30.1.1 BARRY_ZERO_NETWORK

```
#define BARRY_ZERO_NETWORK 0.0
```

Definition at line 85 of file network.hpp.

8.30.1.2 BARRY_ZERO_NETWORK_DENSE

```
#define BARRY_ZERO_NETWORK_DENSE 0
```

Definition at line 86 of file network.hpp.

8.30.1.3 NET_C_DATA_IDX

```
#define NET_C_DATA_IDX(  i \ ) \ ({\tt data.indices[i]})
```

Definition at line 74 of file network.hpp.

8.30.1.4 NET_C_DATA_NUM

```
\label{eq:continuity} \begin{picture}(100,0) \put(0,0){\line(0,0){100}} \put(0,0){\line(0,0){100}}
```

Definition at line 75 of file network.hpp.

8.30.1.5 NETWORK_COUNTER

Value:

```
template<typename Tnet = Network>\
inline double (a) (const Tnet & Array, uint i, uint j, NetCounterData & data)
```

Function for definition of a network counter function

Definition at line 114 of file network.hpp.

8.30.1.6 NETWORK_COUNTER_LAMBDA

Lambda function for definition of a network counter function

Definition at line 119 of file network.hpp.

8.30.1.7 NETWORK_RULE

Function for definition of a network counter function

Definition at line 133 of file network.hpp.

8.30.1.8 NETWORK_RULE_LAMBDA

Lambda function for definition of a network counter function

Definition at line 138 of file network.hpp.

8.30.1.9 NETWORKDENSE_COUNTER_LAMBDA

Definition at line 123 of file network.hpp.

8.30.2 Typedef Documentation

8.30.2.1 NetCounter

```
template<typename Tnet = Network>
using NetCounter = Counter<Tnet, NetCounterData >
```

Definition at line 89 of file network.hpp.

8.30.2.2 NetCounters

```
template<typename Tnet = Network>
using NetCounters = Counters<Tnet, NetCounterData>
```

Definition at line 92 of file network.hpp.

8.30.2.3 NetModel

```
template<typename Tnet >
using NetModel = Model<Tnet, NetCounterData>
```

Definition at line 101 of file network.hpp.

8.30.2.4 NetRule

```
template<typename Tnet = Network>
using NetRule = Rule<Tnet, bool>
```

Definition at line 104 of file network.hpp.

8.30.2.5 NetRules

```
template<typename Tnet = Network>
using NetRules = Rules<Tnet, bool>
```

Definition at line 107 of file network.hpp.

8.30.2.6 NetStatsCounter

```
template<typename Tnet = Network>
using NetStatsCounter = StatsCounter<Tnet, NetCounterData>
```

Definition at line 98 of file network.hpp.

8.30.2.7 NetSupport

```
template<typename Tnet = Network>
using NetSupport = Support<Tnet, NetCounterData >
```

Definition at line 95 of file network.hpp.

8.30.2.8 Network

```
typedef BArray<double, NetworkData> Network
```

Definition at line 82 of file network.hpp.

8.30.2.9 NetworkDense

```
typedef BArrayDense<int, NetworkData> NetworkDense
```

Definition at line 83 of file network.hpp.

8.30.3 Function Documentation

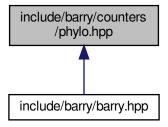
8.30.3.1 rules_zerodiag()

Number of edges.

Definition at line 1383 of file network.hpp.

8.31 include/barry/counters/phylo.hpp File Reference

This graph shows which files directly or indirectly include this file:



Classes

- · class NodeData
 - Data definition for the PhyloArray class.
- · class PhyloCounterData
- class PhyloRuleDynData

Macros

- #define DEFAULT_DUPLICATION 1u
- #define DUPL_SPEC 0u
- #define DUPL_DUPL 1u
- #define DUPL_EITH 2u
- #define MAKE_DUPL_VARS()
- #define IS_EITHER() (DATA_AT == DUPL_EITH)
- #define IS_DUPLICATION() ((DATA_AT == DUPL_DUPL) & (DPL))
- #define IS_SPECIATION() ((DATA_AT == DUPL_SPEC) & (!DPL))
- #define IF_MATCHES()
- #define IF_NOTMATCHES()
- #define PHYLO_COUNTER_LAMBDA(a)

Extension of a simple counter.

- #define PHYLO_RULE_DYN_LAMBDA(a)
- #define PHYLO_CHECK_MISSING()

Typedefs

typedef std::vector< std::pair< uint, uint > > PhyloRuleData

Convenient typedefs for Node objects.

- typedef BArrayDense< uint, NodeData > PhyloArray
- typedef Counter< PhyloArray, PhyloCounterData > PhyloCounter
- typedef Counters
 PhyloArray, PhyloCounterData > PhyloCounters
- typedef Rule < PhyloArray, PhyloRuleData > PhyloRule
- typedef Rules
 PhyloArray, PhyloRuleData > PhyloRules
- typedef Rule< PhyloArray, PhyloRuleDynData > PhyloRuleDyn
- typedef Rules< PhyloArray, PhyloRuleDynData > PhyloRulesDyn
- typedef Support < PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloSupport
- typedef StatsCounter< PhyloArray, PhyloCounterData > PhyloStatsCounter
- typedef Model < PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloModel
- typedef PowerSet
 PhyloArray, PhyloRuleData > PhyloPowerSet

Functions

- std::string get last name (unsigned int d)
- void counter_overall_gains (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)
 Overall functional gains.
- void counter_gains (PhyloCounters *counters, std::vector< uint > nfun, unsigned int duplication=DEFAULT_DUPLICATION)

 Functional gains for a specific function (nfun).
- void counter_gains_k_offspring (PhyloCounters *counters, std::vector< uint > nfun, uint k=1u, unsigned int duplication=DEFAULT_DUPLICATION)

k genes gain function nfun

- void counter_genes_changing (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)
- void counter_preserve_pseudogene (PhyloCounters *counters, unsigned int nfunA, unsigned int nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

Keeps track of how many pairs of genes preserve pseudostate.

- void counter_prop_genes_changing (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)
- void counter_overall_loss (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)
 Overall functional loss.
- void counter_maxfuns (PhyloCounters *counters, uint lb, uint ub, unsigned int duplication=DEFAULT_DUPLICATION)

 Cap the number of functions per gene.
- void counter_loss (PhyloCounters *counters, std::vector< uint > nfun, unsigned int duplication=DEFAULT_DUPLICATION)

 Total count of losses for an specific function.
- void counter_overall_changes (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)
- Total number of changes. Use this statistic to account for "preservation".
- void counter_subfun (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)
 Total count of Sub-functionalization events.
- void counter_cogain (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)
 Co-evolution (joint gain or loss)
- void counter_longest (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)
- Longest branch mutates (either by gain or by loss)
- void counter_neofun (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

 Total number of neofunctionalization events.
- void counter_pairwise_neofun_singlefun (PhyloCounters *counters, uint nfunA, unsigned int duplication=DEFAULT_DUPLICATI

 Total number of neofunctionalization events sum_u sum_{w < u} [x(u,a)*(1 x(w,a)) + (1 x(u,a)) * x(w,a)] change

 stat: delta{x(u,a): 0->1} = 1 2 * x(w,a)
- void counter_neofun_a2b (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION

 Total number of neofunctionalization events.
- void counter_co_opt (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT DUPLICATION)

Function co-opting.

- void counter_k_genes_changing (PhyloCounters *counters, unsigned int k, unsigned int duplication=DEFAULT_DUPLICATION)

 Indicator function. Equals to one if k genes changed and zero otherwise.
- void counter_less_than_p_prop_genes_changing (PhyloCounters *counters, double p, unsigned int duplication=DEFAULT DUPLICATION)

Indicator function. Equals to one if k genes changed and zero otherwise.

- void counter_gains_from_0 (PhyloCounters *counters, std::vector< uint > nfun, unsigned int duplication=DEFAULT_DUPLICAT

 Used when all the functions are in 0 (like the root node prob.)
- void counter_overall_gains_from_0 (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Used when all the functions are in 0 (like the root node prob.)
- void counter_pairwise_overall_change (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Used when all the functions are in 0 (like the root node prob.)
- void counter_pairwise_preserving (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

Used when all the functions are in 0 (like the root node prob.)

 void counter_pairwise_first_gain (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT DUPLICATION)

Used when all the functions are in 0 (like the root node prob.)

void rule_dyn_limit_changes (PhyloSupport *support, uint pos, uint lb, uint ub, unsigned int duplication=DEFAULT_DUPLICATIC
 Overall functional gains.

8.31.1 Macro Definition Documentation

8.31.1.1 DEFAULT_DUPLICATION

#define DEFAULT_DUPLICATION 1u

Definition at line 5 of file phylo.hpp.

8.31.1.2 DUPL_DUPL

#define DUPL_DUPL 1u

Definition at line 7 of file phylo.hpp.

8.31.1.3 DUPL EITH

#define DUPL_EITH 2u

Definition at line 8 of file phylo.hpp.

8.31.1.4 DUPL_SPEC

```
#define DUPL_SPEC Ou
```

Definition at line 6 of file phylo.hpp.

8.31.1.5 IF_MATCHES

```
#define IF_MATCHES()

Value:
    MAKE_DUPL_VARS() \
    if (IS_EITHER() | IS_DUPLICATION() | IS_SPECIATION())
```

Definition at line 19 of file phylo.hpp.

8.31.1.6 IF_NOTMATCHES

```
#define IF_NOTMATCHES( )

Value:
    MAKE_DUPL_VARS() \
    if (!IS_EITHER() & !IS_DUPLICATION() & !IS_SPECIATION())
```

Definition at line 21 of file phylo.hpp.

8.31.1.7 IS_DUPLICATION

```
#define IS_DUPLICATION() ((DATA_AT == DUPL_DUPL) & (DPL))
```

Definition at line 16 of file phylo.hpp.

8.31.1.8 IS_EITHER

```
#define IS_EITHER( ) (DATA_AT == DUPL_EITH)
```

Definition at line 15 of file phylo.hpp.

8.31.1.9 IS_SPECIATION

```
#define IS_SPECIATION( ) ((DATA_AT == DUPL_SPEC) & (!DPL))
```

Definition at line 17 of file phylo.hpp.

8.31.1.10 MAKE_DUPL_VARS

```
#define MAKE_DUPL_VARS( )

Value:
    bool DPL = Array.D_ptr()->duplication; \
```

unsigned int DATA_AT = data[0u];

8.31.1.11 PHYLO_CHECK_MISSING

Definition at line 11 of file phylo.hpp.

```
#define PHYLO_CHECK_MISSING( )

Value:
    if (Array.D_ptr() == nullptr) \
    throw std::logic_error("The array data is nullptr."); \
```

Definition at line 139 of file phylo.hpp.

8.31.1.12 PHYLO_COUNTER_LAMBDA

Extension of a simple counter.

It allows specifying extra arguments, in particular, the corresponding sets of rows to which this statistic may be relevant. This could be important in the case of, for example, counting correlation type statistics between function 1 and 2, and between function 1 and 3.

Definition at line 133 of file phylo.hpp.

8.31.1.13 PHYLO_RULE_DYN_LAMBDA

Value:

```
Rule_fun_type<PhyloArray, PhyloRuleDynData> a = \
[](const PhyloArray & Array, uint i, uint j, PhyloRuleDynData & data)
```

Definition at line 136 of file phylo.hpp.

8.31.2 Typedef Documentation

8.31.2.1 PhyloArray

```
typedef BArrayDense<uint, NodeData> PhyloArray
```

Definition at line 106 of file phylo.hpp.

8.31.2.2 PhyloCounter

```
typedef Counter<PhyloArray, PhyloCounterData > PhyloCounter
```

Definition at line 107 of file phylo.hpp.

8.31.2.3 PhyloCounters

```
typedef Counters< PhyloArray, PhyloCounterData> PhyloCounters
```

Definition at line 108 of file phylo.hpp.

8.31.2.4 PhyloModel

 ${\tt typedef\ Model < PhyloArray,\ PhyloCounterData,\ PhyloRuleData,\ PhyloRuleDynData > PhyloModel}$

Definition at line 118 of file phylo.hpp.

8.31.2.5 PhyloPowerSet

typedef PowerSet<PhyloArray, PhyloRuleData> PhyloPowerSet

Definition at line 119 of file phylo.hpp.

8.31.2.6 PhyloRule

typedef Rule<PhyloArray,PhyloRuleData> PhyloRule

Definition at line 110 of file phylo.hpp.

8.31.2.7 PhyloRuleData

typedef std::vector< std::pair< uint, uint > > PhyloRuleData

Definition at line 99 of file phylo.hpp.

8.31.2.8 PhyloRuleDyn

typedef Rule<PhyloArray,PhyloRuleDynData> PhyloRuleDyn

Definition at line 113 of file phylo.hpp.

8.31.2.9 PhyloRules

typedef Rules<PhyloArray,PhyloRuleData> PhyloRules

Definition at line 111 of file phylo.hpp.

8.31.2.10 PhyloRulesDyn

typedef Rules<PhyloArray,PhyloRuleDynData> PhyloRulesDyn

Definition at line 114 of file phylo.hpp.

8.31.2.11 PhyloStatsCounter

```
typedef StatsCounter<PhyloArray, PhyloCounterData> PhyloStatsCounter
```

Definition at line 117 of file phylo.hpp.

8.31.2.12 PhyloSupport

```
typedef Support<PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloSupport
```

Definition at line 116 of file phylo.hpp.

8.31.3 Function Documentation

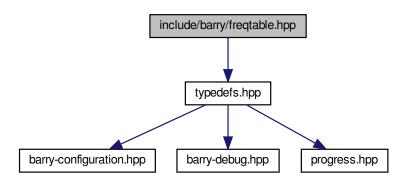
8.31.3.1 get_last_name()

```
std::string get_last_name (
          unsigned int d ) [inline]
```

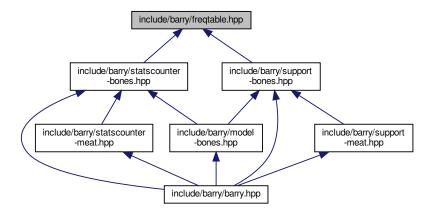
Definition at line 142 of file phylo.hpp.

8.32 include/barry/freqtable.hpp File Reference

```
#include "typedefs.hpp"
Include dependency graph for freqtable.hpp:
```



This graph shows which files directly or indirectly include this file:



Classes

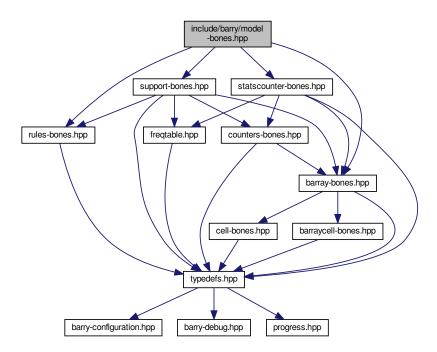
class FreqTable
 T >

Frequency table of vectors.

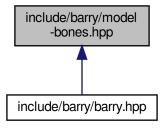
8.33 include/barry/model-bones.hpp File Reference

```
#include "barray-bones.hpp"
#include "support-bones.hpp"
#include "statscounter-bones.hpp"
#include "rules-bones.hpp"
```

Include dependency graph for model-bones.hpp:



This graph shows which files directly or indirectly include this file:



Classes

class Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >

General framework for discrete exponential models. This class allows generating discrete exponential models in the form of a linear exponential model:

Functions

template<typename Array_Type >
 std::vector< double > keygen_default (const Array_Type &Array_)
 Array Hasher class (used for computing support)

8.33.1 Function Documentation

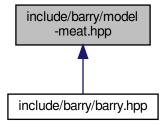
8.33.1.1 keygen_default()

Array Hasher class (used for computing support)

Definition at line 16 of file model-bones.hpp.

8.34 include/barry/model-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



Macros

- #define MODEL_TYPE()
- #define MODEL TEMPLATE ARGS()
- #define MODEL_TEMPLATE(a, b) template MODEL_TEMPLATE_ARGS() inline a MODEL_TYPE()::b

Functions

- double update_normalizing_constant (const double *params, const double *support, size_t k, size_t n)
- double likelihood_ (const double *stats_target, const std::vector< double > ¶ms, const double normalizing_constant, size_t n_params, bool log_=false)
- MODEL_TEMPLATE (, Model)()
- MODEL_TEMPLATE (, Model)(const MODEL_TYPE() &Model_)

8.34.1 Macro Definition Documentation

8.34.1.1 MODEL_TEMPLATE

Definition at line 89 of file model-meat.hpp.

8.34.1.2 MODEL_TEMPLATE_ARGS

Definition at line 86 of file model-meat.hpp.

8.34.1.3 MODEL_TYPE

Definition at line 83 of file model-meat.hpp.

8.34.2 Function Documentation

8.34.2.1 likelihood_()

Definition at line 47 of file model-meat.hpp.

8.34.2.2 MODEL_TEMPLATE() [1/2]

```
MODEL_TEMPLATE (

Model )
```

Definition at line 93 of file model-meat.hpp.

8.34.2.3 MODEL_TEMPLATE() [2/2]

```
MODEL_TEMPLATE (

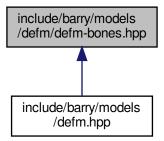
Model ) const &
```

Definition at line 157 of file model-meat.hpp.

8.34.2.4 update_normalizing_constant()

Definition at line 11 of file model-meat.hpp.

8.35 include/barry/models/defm/defm-bones.hpp File Reference



Classes

class Geese

Annotated Phylo Model.

Macros

• #define INITIALIZED()

Functions

- RULE_FUNCTION (rule_empty_free)
- std::vector< double > keygen_full (const phylocounters::PhyloArray &array)
- bool vec diff (const std::vector< unsigned int > &s, const std::vector< unsigned int > &a)

8.35.1 Macro Definition Documentation

8.35.1.1 INITIALIZED

```
#define INITIALIZED( )

Value:
    if (!this->initialized) \
        throw std::logic_error("The model has not been initialized yet.");
```

Definition at line 9 of file defm-bones.hpp.

8.35.2 Function Documentation

8.35.2.1 keygen_full()

Definition at line 20 of file defm-bones.hpp.

8.35.2.2 RULE_FUNCTION()

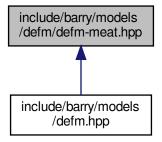
Definition at line 13 of file defm-bones.hpp.

8.35.2.3 vec_diff()

Definition at line 44 of file defm-bones.hpp.

8.36 include/barry/models/defm/defm-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



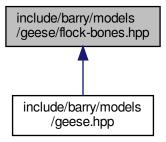
8.37 include/barry/models/geese.hpp File Reference

```
#include "geese/geese-node-bones.hpp"
#include "geese/geese-bones.hpp"
#include "geese/geese-meat.hpp"
#include "geese/geese-meat-constructors.hpp"
#include "geese/geese-meat-likelihood.hpp"
#include "geese/geese-meat-likelihood_exhaust.hpp"
#include "geese/geese-meat-simulate.hpp"
#include "geese/geese-meat-predict.hpp"
#include "geese/geese-meat-predict_exhaust.hpp"
#include "geese/geese-meat-predict_sim.hpp"
#include "geese/flock-bones.hpp"
#include "geese/flock-meat.hpp"
Include dependency graph for geese.hpp:
```



8.38 include/barry/models/geese/flock-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:

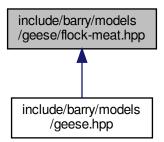


Classes

· class Flock

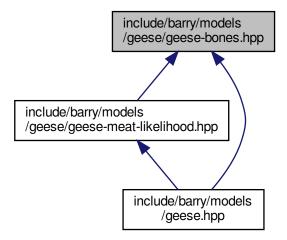
A Flock is a group of Geese.

8.39 include/barry/models/geese/flock-meat.hpp File Reference



8.40 include/barry/models/geese/geese-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



Classes

• class Geese

Annotated Phylo Model.

Macros

• #define INITIALIZED()

Functions

- template<typename Ta , typename Tb > $std::vector < Ta > vector_caster \ (const \ std::vector < Tb > \&x)$
- RULE_FUNCTION (rule_empty_free)
- std::vector< double > keygen_full (const phylocounters::PhyloArray &array)
- bool vec_diff (const std::vector< unsigned int > &s, const std::vector< unsigned int > &a)

8.40.1 Macro Definition Documentation

8.40.1.1 INITIALIZED

```
#define INITIALIZED()
Value:
    if (!this->initialized) \
    throw std::logic_error("The model has not been initialized yet.");
```

Definition at line 22 of file geese-bones.hpp.

8.40.2 Function Documentation

8.40.2.1 keygen_full()

Definition at line 35 of file geese-bones.hpp.

8.40.2.2 RULE_FUNCTION()

Definition at line 26 of file geese-bones.hpp.

8.40.2.3 vec_diff()

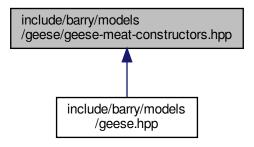
Definition at line 59 of file geese-bones.hpp.

8.40.2.4 vector_caster()

Definition at line 10 of file geese-bones.hpp.

8.41 include/barry/models/geese/geese-meat-constructors.hpp File Reference

This graph shows which files directly or indirectly include this file:

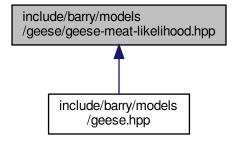


8.42 include/barry/models/geese/geese-meat-likelihood.hpp File Reference

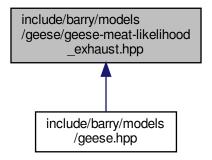
#include "geese-bones.hpp"
Include dependency graph for geese-meat-likelihood.hpp:



This graph shows which files directly or indirectly include this file:

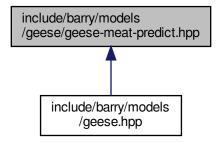


8.43 include/barry/models/geese/geese-meat-likelihood_exhaust.hpp File Reference

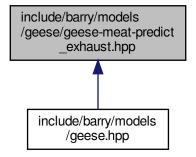


8.44 include/barry/models/geese/geese-meat-predict.hpp File Reference

This graph shows which files directly or indirectly include this file:



8.45 include/barry/models/geese/geese-meat-predict_exhaust.hpp File Reference



8.46 include/barry/models/geese/geese-meat-predict_sim.hpp File Reference

This graph shows which files directly or indirectly include this file:



8.47 include/barry/models/geese/geese-meat-simulate.hpp File Reference



8.48 include/barry/models/geese/geese-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



8.49 include/barry/models/geese/geese-node-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



Classes

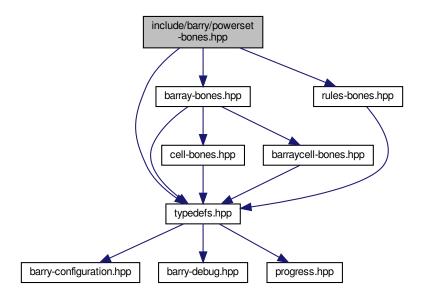
• class Node

A single node for the model.

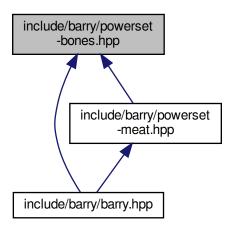
8.50 include/barry/powerset-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "barray-bones.hpp"
#include "rules-bones.hpp"
```

Include dependency graph for powerset-bones.hpp:



This graph shows which files directly or indirectly include this file:



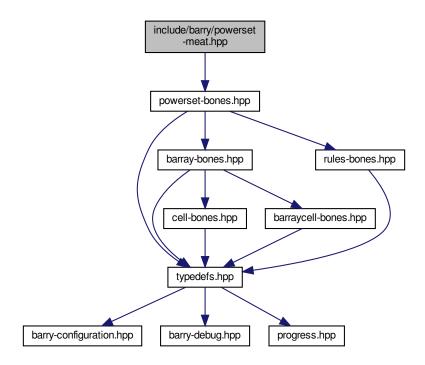
Classes

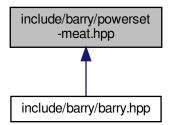
- class PowerSet < Array_Type, Data_Rule_Type >

Powerset of a binary array.

8.51 include/barry/powerset-meat.hpp File Reference

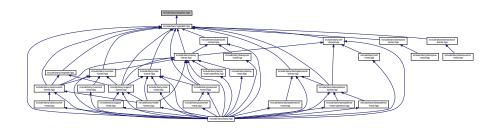
#include "powerset-bones.hpp"
Include dependency graph for powerset-meat.hpp:





8.52 include/barry/progress.hpp File Reference

This graph shows which files directly or indirectly include this file:



Classes

class Progress

A simple progress bar.

Macros

• #define BARRY_PROGRESS_BAR_WIDTH 80

8.52.1 Macro Definition Documentation

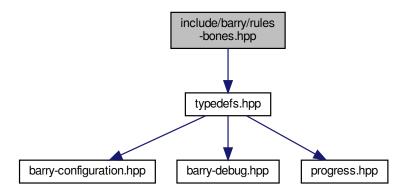
8.52.1.1 BARRY_PROGRESS_BAR_WIDTH

#define BARRY_PROGRESS_BAR_WIDTH 80

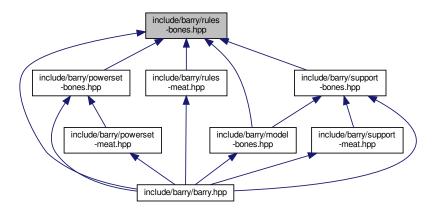
Definition at line 5 of file progress.hpp.

8.53 include/barry/rules-bones.hpp File Reference

#include "typedefs.hpp"
Include dependency graph for rules-bones.hpp:



This graph shows which files directly or indirectly include this file:



Classes

- class Rule < Array_Type, Data_Type >
 Rule for determining if a cell should be included in a sequence.
- class Rules < Array_Type, Data_Type >
 Vector of objects of class Rule.

Functions

template < typename Array_Type , typename Data_Type >
 bool rule_fun_default (const Array_Type *array, uint i, uint j, Data_Type *dat)

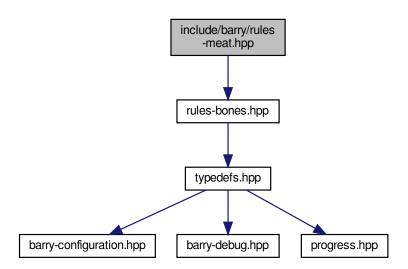
8.53.1 Function Documentation

8.53.1.1 rule_fun_default()

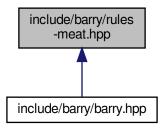
Definition at line 7 of file rules-bones.hpp.

8.54 include/barry/rules-meat.hpp File Reference

#include "rules-bones.hpp"
Include dependency graph for rules-meat.hpp:



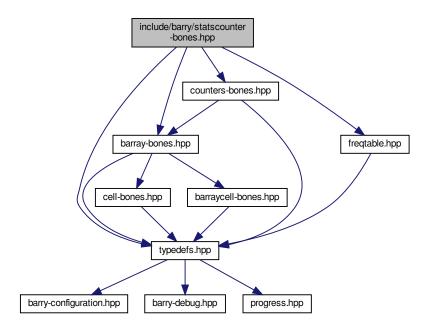
This graph shows which files directly or indirectly include this file:



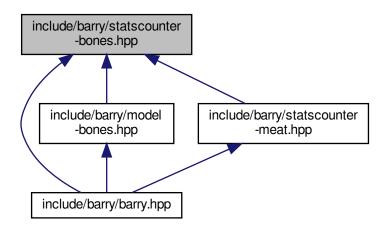
8.55 include/barry/statscounter-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "barray-bones.hpp"
#include "freqtable.hpp"
```

#include "counters-bones.hpp"
Include dependency graph for statscounter-bones.hpp:



This graph shows which files directly or indirectly include this file:

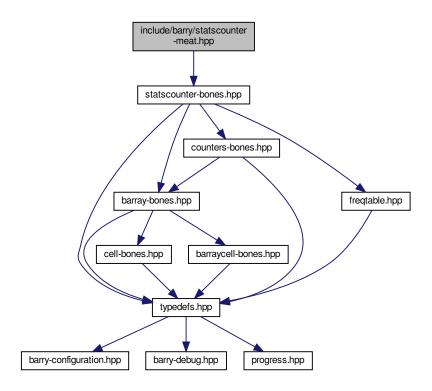


Classes

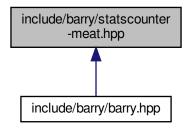
class StatsCounter < Array_Type, Data_Type >
 Count stats for a single Array.

8.56 include/barry/statscounter-meat.hpp File Reference

#include "statscounter-bones.hpp"
Include dependency graph for statscounter-meat.hpp:



This graph shows which files directly or indirectly include this file:



Macros

- #define STATSCOUNTER_TYPE() StatsCounter<Array_Type,Data_Type>
- #define STATSCOUNTER_TEMPLATE_ARGS() < typename Array_Type, typename Data_Type>
- #define STATSCOUNTER_TEMPLATE(a, b) template STATSCOUNTER_TEMPLATE_ARGS() inline a STATSCOUNTER_TYPE()::b

Functions

- STATSCOUNTER_TEMPLATE (, StatsCounter)(const StatsCounter< Array_Type
- EmptyArray clear ()
- STATSCOUNTER_TEMPLATE (,~StatsCounter)()
- STATSCOUNTER_TEMPLATE (void, reset_array)(const Array_Type *Array_)
- STATSCOUNTER_TEMPLATE (void, add_counter)(Counter< Array_Type
- STATSCOUNTER_TEMPLATE (void, set_counters)(Counters < Array_Type
- STATSCOUNTER_TEMPLATE (void, count_init)(uint i
- current stats resize (counters->size(), 0.0)
- for (uint n=0u;n< counters->size();++n) current stats[n]
- STATSCOUNTER_TEMPLATE (void, count_current)(uint i
- STATSCOUNTER_TEMPLATE (std::vector< std::string >, get_names)() const
- STATSCOUNTER_TEMPLATE (std::vector< std::string >, get_descriptions)() const

Variables

- · Data_Type & counter
- EmptyArray = *Array
- current stats = counter.current stats
- counters = new Counters
 Array_Type,Data_Type>((*counter.counters))
- counter_deleted = false
- Data_Type f_
- return
- Data_Type * counters_
- uint j

8.56.1 Macro Definition Documentation

8.56.1.1 STATSCOUNTER_TEMPLATE

Definition at line 11 of file statscounter-meat.hpp.

8.56.1.2 STATSCOUNTER TEMPLATE ARGS

```
template STATSCOUNTER_TEMPLATE_ARGS() <typename Array_Type</pre>, typename Data_Type>
```

Definition at line 9 of file statscounter-meat.hpp.

8.56.1.3 STATSCOUNTER_TYPE

```
template Data_Type * STATSCOUNTER_TYPE( ) StatsCounter<Array_Type,Data_Type>
```

Definition at line 7 of file statscounter-meat.hpp.

8.56.2 Function Documentation

8.56.2.1 clear()

```
EmptyArray clear ( )
```

8.56.2.2 for()

8.56.2.3 resize()

8.56.2.4 STATSCOUNTER_TEMPLATE() [1/9]

```
STATSCOUNTER_TEMPLATE (
StatsCounter ) const
```

8.56.2.5 STATSCOUNTER_TEMPLATE() [2/9]

```
STATSCOUNTER_TEMPLATE ( \sim StatsCounter )
```

Definition at line 30 of file statscounter-meat.hpp.

8.56.2.6 STATSCOUNTER_TEMPLATE() [3/9]

```
STATSCOUNTER_TEMPLATE (
          std::vector< std::string > ,
          get_descriptions ) const
```

Definition at line 259 of file statscounter-meat.hpp.

8.56.2.7 STATSCOUNTER_TEMPLATE() [4/9]

Definition at line 254 of file statscounter-meat.hpp.

8.56.2.8 STATSCOUNTER_TEMPLATE() [5/9]

8.56.2.9 STATSCOUNTER_TEMPLATE() [6/9]

8.56.2.10 STATSCOUNTER_TEMPLATE() [7/9]

8.56.2.11 STATSCOUNTER_TEMPLATE() [8/9]

Definition at line 37 of file statscounter-meat.hpp.

8.56.2.12 STATSCOUNTER_TEMPLATE() [9/9]

8.56.3 Variable Documentation

8.56.3.1 counter

```
Data_Type& counter

Initial value:
{
    Array = counter.Array
```

Definition at line 15 of file statscounter-meat.hpp.

8.56.3.2 counter_deleted

```
counter_deleted = false
```

Definition at line 26 of file statscounter-meat.hpp.

8.56.3.3 counters

```
counters = new Counters<Array_Type, Data_Type>((*counter.counters))
```

Definition at line 25 of file statscounter-meat.hpp.

8.56.3.4 counters_

```
Data_Type* counters_
Initial value:
{
    if (!counter_deleted)
```

Definition at line 56 of file statscounter-meat.hpp.

8.56.3.5 current_stats

```
current_stats = counter.current_stats
```

Definition at line 22 of file statscounter-meat.hpp.

8.56.3.6 EmptyArray

```
EmptyArray = *Array
```

Definition at line 20 of file statscounter-meat.hpp.

8.56.3.7 f_

```
Data_Rule_Dyn_Type f_
```

Initial value:

```
counters->add_counter(f_)
```

Definition at line 47 of file statscounter-meat.hpp.

8.56.3.8 j

```
uint j
```

Initial value:

```
if (counters->size() == 0u)
    throw std::logic_error("No counters added: Cannot count without knowning what to count!")
```

Definition at line 69 of file statscounter-meat.hpp.

8.56.3.9 return

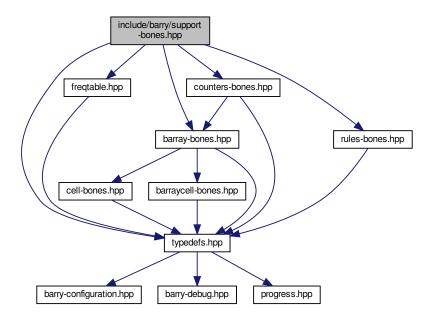
return

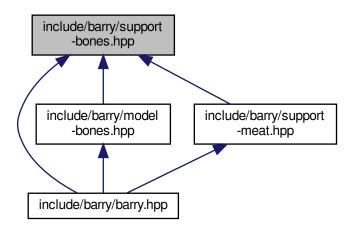
Definition at line 52 of file statscounter-meat.hpp.

8.57 include/barry/support-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "barray-bones.hpp"
#include "freqtable.hpp"
#include "counters-bones.hpp"
#include "rules-bones.hpp"
```

Include dependency graph for support-bones.hpp:



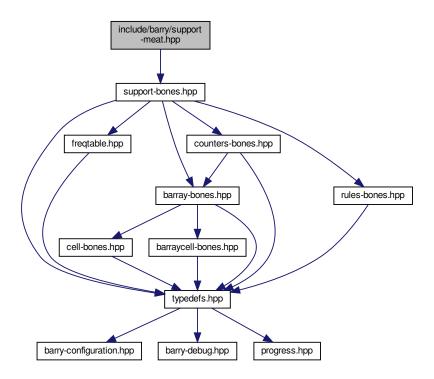


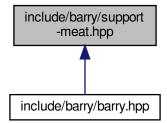
Classes

class Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >
 Compute the support of sufficient statistics.

8.58 include/barry/support-meat.hpp File Reference

#include "support-bones.hpp"
Include dependency graph for support-meat.hpp:





Macros

- #define BARRY_SUPPORT_MEAT_HPP 1
- #define SUPPORT TEMPLATE ARGS()
- #define SUPPORT TYPE()
- #define SUPPORT_TEMPLATE(a, b)

Functions

- SUPPORT_TEMPLATE (void, init_support)(std
- SUPPORT TEMPLATE (void, reset array)()
- SUPPORT_TEMPLATE (void, reset_array)(const Array_Type &Array_)
- SUPPORT TEMPLATE (void, calc backend sparse)(uint pos
- calc_backend_sparse (pos+1u, array_bank, stats_bank)
- EmptyArray insert_cell (coord_i, coord_j, EmptyArray.default_val().value, false, false)
- for (uint n=0u;n< n_counters;++n)
- if (rules dyn->size() > 0u)
- if (array_bank !=nullptr) array_bank -> push_back(EmptyArray)
- EmptyArray rm cell (coord i, coord j, false, false)
- if (change_stats_different > 0u)
- SUPPORT TEMPLATE (void, calc backend dense)(uint pos
- calc_backend_dense (pos+1u, array_bank, stats_bank)
- EmptyArray insert cell (coord i, coord j, 1, false, false)
- SUPPORT_TEMPLATE (void, calc)(std
- SUPPORT TEMPLATE (void, add counter)(Counter< Array Type
- SUPPORT_TEMPLATE (void, set_counters)(Counters< Array_Type
- SUPPORT TEMPLATE (void, add rule)(Rule< Array Type
- SUPPORT TEMPLATE (void, set rules)(Rules < Array Type
- SUPPORT_TEMPLATE (void, add_rule_dyn)(Rule < Array_Type
- SUPPORT_TEMPLATE (void, set_rules_dyn)(Rules< Array_Type
- SUPPORT_TEMPLATE (bool, eval_rules_dyn)(const std
- SUPPORT_TEMPLATE (std::vector< double >, get_counts)() const
- SUPPORT_TEMPLATE (std::vector< double > *, get_current_stats)()
- SUPPORT_TEMPLATE (void, print)() const
- SUPPORT_TEMPLATE (const FreqTable<> &, get_data)() const

Variables

- std::vector< Array_Type > * array_bank
- std::vector< Array Type > std::vector< double > * stats bank
- const size t & coord i = coordinates free[pos * 2u]
- const size_t & coord_j = coordinates_free[pos * 2u + 1u]
- · double tmp_chng
- unsigned int change stats different = hashes initialized[pos] ? Ou : 1u
- else
- · & hashes [pos]
- return
- Data Counter Type f
- Data_Counter_Type * counters_
- delete_counters = false
- counters = counters
- Data_Rule_Type * rules_
- delete rules = false
- rules = rules
- delete_rules_dyn = false
- rules_dyn = rules_

8.58.1 Macro Definition Documentation

8.58.1.1 BARRY_SUPPORT_MEAT_HPP

```
#define BARRY_SUPPORT_MEAT_HPP 1
```

Definition at line 4 of file support-meat.hpp.

8.58.1.2 SUPPORT_TEMPLATE

Value:

```
template SUPPORT_TEMPLATE_ARGS() \
inline a SUPPORT_TYPE()::b
```

Definition at line 12 of file support-meat.hpp.

8.58.1.3 SUPPORT_TEMPLATE_ARGS

```
template SUPPORT_TEMPLATE_ARGS( )
```

Value:

```
<tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre><tpre>
```

Definition at line 6 of file support-meat.hpp.

8.58.1.4 SUPPORT_TYPE

```
template Data_Rule_Dyn_Type * SUPPORT_TYPE( )
```

Value

```
Support<Array_Type, Data_Counter_Type, Data_Rule_Type, \
Data_Rule_Dyn_Type>
```

Definition at line 9 of file support-meat.hpp.

8.58.2 Function Documentation

8.58.2.1 calc_backend_dense()

```
calc_backend_dense (
    pos+ 1u,
    array_bank ,
    stats_bank )
```

8.58.2.2 calc_backend_sparse()

```
calc_backend_sparse ( pos+\ 1u, array\_bank\ , stats\_bank\ )
```

8.58.2.3 for()

```
for ( )
```

Definition at line 162 of file support-meat.hpp.

```
8.58.2.4 if() [1/3]
```

```
if (
     array_bank ! = nullptr ) -> push_back(EmptyArray)
```

8.58.2.5 if() [2/3]

```
if ( \label{eq:change_stats_different} \mbox{,} \\ \mbox{Ou )}
```

Definition at line 242 of file support-meat.hpp.

8.58.2.6 if() [3/3]

```
if (
    rules_dyn-> size(),
    0u )
```

Definition at line 190 of file support-meat.hpp.

8.58.2.7 insert_cell() [1/2]

8.58.2.8 insert_cell() [2/2]

8.58.2.9 rm_cell()

8.58.2.10 SUPPORT_TEMPLATE() [1/17]

```
SUPPORT_TEMPLATE (
          bool ,
          eval_rules_dyn ) const
```

Definition at line 491 of file support-meat.hpp.

8.58.2.11 SUPPORT_TEMPLATE() [2/17]

Definition at line 560 of file support-meat.hpp.

8.58.2.12 SUPPORT_TEMPLATE() [3/17]

```
SUPPORT_TEMPLATE (
          std::vector< double > * ,
          get_current_stats )
```

Definition at line 545 of file support-meat.hpp.

8.58.2.13 SUPPORT_TEMPLATE() [4/17]

```
SUPPORT_TEMPLATE (
          std::vector< double > ,
          get_counts ) const
```

Definition at line 533 of file support-meat.hpp.

8.58.2.14 SUPPORT_TEMPLATE() [5/17]

8.58.2.15 SUPPORT_TEMPLATE() [6/17]

8.58.2.16 SUPPORT_TEMPLATE() [7/17]

8.58.2.17 SUPPORT_TEMPLATE() [8/17]

```
SUPPORT_TEMPLATE (
     void ,
     calc )
```

Definition at line 374 of file support-meat.hpp.

8.58.2.18 SUPPORT_TEMPLATE() [9/17]

8.58.2.19 SUPPORT_TEMPLATE() [10/17]

8.58.2.20 SUPPORT_TEMPLATE() [11/17]

Definition at line 16 of file support-meat.hpp.

8.58.2.21 SUPPORT_TEMPLATE() [12/17]

```
SUPPORT_TEMPLATE (
     void ,
     print ) const
```

Definition at line 549 of file support-meat.hpp.

8.58.2.22 SUPPORT_TEMPLATE() [13/17]

```
SUPPORT_TEMPLATE (
     void ,
     reset_array )
```

Definition at line 117 of file support-meat.hpp.

8.58.2.23 SUPPORT_TEMPLATE() [14/17]

Definition at line 123 of file support-meat.hpp.

8.58.2.24 SUPPORT_TEMPLATE() [15/17]

8.58.2.25 SUPPORT_TEMPLATE() [16/17]

```
SUPPORT_TEMPLATE (
     void ,
     set_rules )
```

8.58.2.26 SUPPORT_TEMPLATE() [17/17]

8.58.3 Variable Documentation

8.58.3.1 array_bank

```
std::vector< Array_Type > * array_bank
```

Definition at line 134 of file support-meat.hpp.

8.58.3.2 change_stats_different

```
unsigned int change_stats_different = hashes_initialized[pos] ? Ou : 1u
```

Definition at line 161 of file support-meat.hpp.

8.58.3.3 coord_i

```
const size_t & coord_i = coordinates_free[pos * 2u]
```

Definition at line 147 of file support-meat.hpp.

8.58.3.4 coord_j

```
const size_t & coord_j = coordinates_free[pos * 2u + 1u]
```

Definition at line 148 of file support-meat.hpp.

8.58.3.5 counters

```
counters = counters_
```

Definition at line 419 of file support-meat.hpp.

8.58.3.6 counters_

```
Data_Counter_Type* counters_
Initial value:
{
```

if (delete_counters)
 delete counters

Definition at line 412 of file support-meat.hpp.

8.58.3.7 delete_counters

```
delete_counters = false
```

Definition at line 418 of file support-meat.hpp.

8.58.3.8 delete_rules

```
delete_rules = false
```

Definition at line 452 of file support-meat.hpp.

8.58.3.9 delete_rules_dyn

```
delete_rules_dyn = false
```

Definition at line 484 of file support-meat.hpp.

8.58.3.10 else

Definition at line 215 of file support-meat.hpp.

8.58.3.11 f_

```
Data_Rule_Dyn_Type f_
Initial value:
{
    counters->add_counter(f_)
```

Definition at line 403 of file support-meat.hpp.

8.58.3.12 hashes

```
& hashes
```

Definition at line 220 of file support-meat.hpp.

8.58.3.13 return

return

Definition at line 254 of file support-meat.hpp.

8.58.3.14 rules

```
rules = rules_
```

Definition at line 453 of file support-meat.hpp.

8.58.3.15 rules_

```
Data_Rule_Dyn_Type * rules_
Initial value:
{
    if (delete_rules)
        delete rules
```

Definition at line 446 of file support-meat.hpp.

8.58.3.16 rules_dyn

```
rules_dyn = rules_
```

Definition at line 485 of file support-meat.hpp.

8.58.3.17 stats_bank

```
std::vector< Array_Type > std::vector< double > * stats_bank

Initial value:
{
    if (pos >= coordiantes_n_free)
        return
```

Definition at line 135 of file support-meat.hpp.

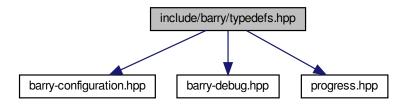
8.58.3.18 tmp_chng

double tmp_chng

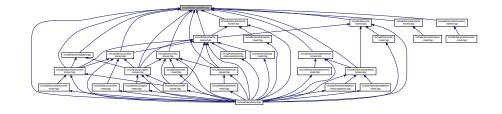
Definition at line 160 of file support-meat.hpp.

8.59 include/barry/typedefs.hpp File Reference

```
#include "barry-configuration.hpp"
#include "barry-debug.hpp"
#include "progress.hpp"
Include dependency graph for typedefs.hpp:
```



This graph shows which files directly or indirectly include this file:



Classes

- class Entries < Cell_Type >
 - A wrapper class to store source, target, val from a BArray object.
- struct vecHasher< T >

Namespaces

• CHECK

Integer constants used to specify which cell should be check.

• EXISTS

Integer constants used to specify which cell should be check to exist or not.

Typedefs

```
typedef unsigned int uint
typedef std::vector< std::pair< std::vector< double >, uint > > Counts_type
template<typename Cell_Type >
using Row_type = Map< uint, Cell< Cell_Type > >
vsing Col_type = Map< uint, Cell< Cell_Type > * >
using Col_type = Map< uint, Cell< Cell_Type > * >
template<typename Ta = double, typename Tb = uint>
using MapVec_type = std::unordered_map< std::vector< Ta >, Tb, vecHasher< Ta > >
template<typename Array_Type , typename Data_Type >
using Counter_fun_type = std::function< double(const Array_Type &, uint, uint, Data_Type &)>
Counter and rule functions.
template<typename Array_Type , typename Data_Type >
```

using Rule_fun_type = std::function< bool(const Array_Type &, uint, uint, Data_Type &)>

Functions

```
    template < typename T >
        T vec_inner_prod (const T *a, const T *b, size_t n)
    template <> double vec_inner_prod (const double *a, const double *b, size_t n)
    template < typename T >
        bool vec_equal (const std::vector < T > &a, const std::vector < T > &b)
        Compares if -a- and -b- are equal.
    template < typename T >
        bool vec_equal_approx (const std::vector < T > &a, const std::vector < T > &b, double eps=1e-100)
```

Variables

```
const int CHECK::BOTH = -1
const int CHECK::NONE = 0
const int CHECK::ONE = 1
const int CHECK::TWO = 2
const int EXISTS::BOTH = -1
const int EXISTS::NONE = 0
const int EXISTS::ONE = 1
const int EXISTS::TWO = 1
const int EXISTS::UKNOWN = -1
const int EXISTS::AS_ZERO = 0
const int EXISTS::AS ONE = 1
```

8.59.1 Typedef Documentation

358 File Documentation

8.59.1.1 Col_type

```
template<typename Cell_Type >
using Col_type = Map< uint, Cell<Cell_Type>* >
```

Definition at line 71 of file typedefs.hpp.

8.59.1.2 Counter_fun_type

```
template<typename Array_Type , typename Data_Type >
using Counter_fun_type = std::function<double(const Array_Type &, uint, uint, Data_Type &)>
```

Counter and rule functions.

Parameters

Array_Type	a BArray
unit,uint	Focal cell
Data_Type	Data associated with the function, for example, id of the attribute in the Array.

Returns

```
Counter_fun_type a double (the change statistic)
Rule_fun_type a bool. True if the cell is blocked.
```

Definition at line 148 of file typedefs.hpp.

8.59.1.3 Counts_type

```
typedef std::vector< std::pair< std::vector<double>, uint > > Counts_type
```

Definition at line 52 of file typedefs.hpp.

8.59.1.4 MapVec_type

```
template<typename Ta = double, typename Tb = uint>
using MapVec_type = std::unordered_map< std::vector< Ta >, Tb, vecHasher<Ta> >
```

Definition at line 129 of file typedefs.hpp.

8.59.1.5 Row_type

```
template<typename Cell_Type >
using Row_type = Map< uint, Cell<Cell_Type> >
```

Definition at line 68 of file typedefs.hpp.

8.59.1.6 Rule_fun_type

```
template<typename Array_Type , typename Data_Type >
using Rule_fun_type = std::function<bool(const Array_Type &, uint, uint, Data_Type &)>
```

Definition at line 151 of file typedefs.hpp.

8.59.1.7 uint

```
typedef unsigned int uint
```

Definition at line 18 of file typedefs.hpp.

8.59.2 Function Documentation

8.59.2.1 vec_equal()

Compares if -a- and -b- are equal.

Parameters

```
a,b Two vectors of the same length
```

Returns

true if all elements are equal.

Definition at line 162 of file typedefs.hpp.

360 File Documentation

8.59.2.2 vec_equal_approx()

Definition at line 180 of file typedefs.hpp.

8.59.2.3 vec_inner_prod() [1/2]

Definition at line 226 of file typedefs.hpp.

8.59.2.4 vec_inner_prod() [2/2]

Definition at line 203 of file typedefs.hpp.

8.60 README.md File Reference

Index

```
\simBArray
                                                      \simPhyloRuleDynData
    BArray< Cell Type, Data Type >, 39
                                                           PhyloRuleDynData, 176
                                                      \simPowerSet
\simBArrayCell
                                                           PowerSet < Array_Type, Data_Rule_Type >, 179
    BArrayCell< Cell_Type, Data_Type >, 50
~BArrayCell const
                                                      \simProgress
    BArrayCell_const< Cell_Type, Data_Type >, 52
                                                           Progress, 184
{\sim}\mathsf{BArrayDense}
                                                      \simRule
    BArrayDense < Cell_Type, Data_Type >, 58
                                                          Rule < Array_Type, Data_Type >, 186
                                                      \simRules
\simBArrayDenseCell
    BArrayDenseCell< Cell_Type, Data_Type >, 70
                                                           Rules < Array_Type, Data_Type >, 188
\simBArrayRow
                                                      \simStatsCounter
    BArrayRow< Cell Type, Data Type >, 83
                                                          StatsCounter< Array Type, Data Type >, 192
~BArrayRow const
                                                      \simSupport
    BArrayRow const< Cell Type, Data Type >, 85
                                                                                      Data Counter Type,
                                                          Support<
                                                                       Array Type,
\simBArrayVector
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
    BArrayVector< Cell Type, Data Type >, 88
~BArrayVector const
                                                      active
    BArrayVector_const< Cell_Type, Data_Type >, 91
                                                           Cell< Cell_Type >, 98
\simCell
                                                      add
    Cell < Cell Type >, 95
                                                          barray-meat.hpp, 226
\simConstBArrayRowIter
                                                          barraydense-meat.hpp, 254
    ConstBArrayRowlter< Cell Type, Data Type >,
                                                          Cell < Cell Type >, 96, 97
                                                          FreqTable< T >, 126
\simCounter
                                                      add array
    Counter< Array_Type, Data_Type >, 103
                                                          Model<
                                                                      Array Type,
                                                                                       Data Counter Type,
\simCounters
                                                               Data Rule Type, Data Rule Dyn Type >,
    Counters < Array_Type, Data_Type >, 107
                                                               151
\simDEFMCounterData
                                                      add_counter
    DEFMCounterData, 111
                                                          Counters < Array_Type, Data_Type >, 108
\simDEFMData
                                                                      Array_Type,
                                                           Model<
                                                                                      Data Counter Type,
    DEFMData, 114
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
\simEntries
                                                               152
    Entries < Cell Type >, 116
                                                          StatsCounter< Array Type, Data Type >, 192
\simFlock
                                                                                      Data_Counter_Type,
                                                           Support<
                                                                       Array Type,
     Flock, 119
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
\simFregTable
                                                               197
    FreqTable < T >, 125
                                                      add data
\simGeese
                                                           Flock, 119
    Geese, 132, 133
                                                      add rule
\simModel
                                                          Model<
                                                                      Array Type,
                                                                                      Data Counter Type,
                Array_Type,
                                Data_Counter_Type,
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
         Data_Rule_Type, Data_Rule_Dyn_Type >,
                                                               152
         151
                                                          PowerSet< Array Type, Data Rule Type >, 179,
\simNetCounterData
                                                               180
    NetCounterData, 162
                                                          Rules < Array Type, Data Type >, 189
\simNetworkData
                                                                       Array_Type,
                                                          Support<
                                                                                      Data Counter Type,
    NetworkData, 165
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
\simNode
                                                               197, 198
    Node, 167
                                                      add_rule_dyn
```

Model< Array_Type, Data_Counter_Type,	operator/=, 45
Data_Rule_Type, Data_Rule_Dyn_Type >,	operator=, 45
152	operator==, 45
Support< Array_Type, Data_Counter_Type,	out_of_range, 45
Data_Rule_Type, Data_Rule_Dyn_Type >,	print, 45
198	reserve, 46
annotations	resize, 46
Node, 168	rm_cell, 46
ans	row, 46
barray-meat.hpp, 217, 226	set_data, 46
barraydense-meat.hpp, 243, 254	swap_cells, 47
Array	swap_cols, 47
ConstBArrayRowlter< Cell_Type, Data_Type >,	swap_rows, 47
100	toggle_cell, 47
array	toggle_lock, 47
Node, 168	transpose, 48
Array_	visited, 49
barray-meat.hpp, 226	zero_col, 48
array_bank	zero_row, 48
support-meat.hpp, 352	barray-bones.hpp
arrays	BARRAY BONES HPP, 208
Node, 169	barray-meat-operators.hpp
AS_ONE	BARRAY_TEMPLATE, 210–212
EXISTS, 33	BARRAY_TEMPLATE_ARGS, 210, 212
as_vector	BARRAY_TYPE, 210, 212
FreqTable < T >, 126	BARRY_BARRAY_MEAT_OPERATORS_HPP,
AS_ZERO	210
	COL, 211
EXISTS, 33	
at DEEMD-11 111	for, 212
DEFMData, 114	operator(), 213
PhyloCounterData, 173	rhs, 213
BArray	ROW, 211
-	this, 213
BArray Cell_Type, Data_Type >, 38, 39	barray-meat.hpp
BArray Cell_Type, Data_Type >, 35	add, 226
~BArray, 39	ans, 217, 226
BArray, 38, 39	Array_, 226
BArrayCell < Cell_Type, Data_Type >, 48	BARRAY_TEMPLATE, 216-221
BArrayCell_const< Cell_Type, Data_Type >, 48	BARRAY_TEMPLATE_ARGS, 216
clear, 39	BARRAY_TYPE, 216
col, 39	check_bounds, 226
D, 40	check exists, 227
D_ptr, 40	COL, 217, 221
default_val, 40	col0, 227
flush_data, 40	const, 227
get_cell, 40	copy_data, 227
get_col_vec, 41	data, 228
get_entries, 41	delete_data, 228
get_row_vec, 41	
insert_cell, 42	delete_data_, 228
is_dense, 42	else, 228
is_empty, 42	false, 228
ncol, 43	first, 229
nnozero, 43	for, 222
	i1, 229
nrow, 43	if, 222–225
operator() 43	j, 229
operator(), 43	
	j0, 229
operator+=, 44	j0, 229 j1, 229
operator-=, 44	

M, 225, 229 M_, 230 N, 230	BArrayDenseCol < Cell_Type, Data_Type >, 68, 75 BArrayDenseCol_const < Cell_Type, Data_Type >, 68
NCells, 230 report, 230	BArrayDenseRow< Cell_Type, Data_Type >, 69,
resize, 225	BArrayDenseRow_const< Cell_Type, Data_Type
return, 225, 230	>, 69
ROW, 217, 225, 226	clear, 58
row0, 231	col, 58, 59
search, 231	colsum, 59
source, 231	D, 59
target, 231	D_ptr, 59
v, 231	default_val, 60
value, 231	get_cell, 60
BARRAY_BONES_HPP	get_col_vec, 60
barray-bones.hpp, 208	get_data, 60
BARRAY_TEMPLATE	get_entries, 60
barray-meat-operators.hpp, 210–212	get_row_vec, 61
barray-meat.hpp, 216–221	insert_cell, 61
BARRAY_TEMPLATE_ARGS	is_dense, 62
barray-meat-operators.hpp, 210, 212	is_empty, 62
barray-meat.hpp, 216	ncol, 62
BARRAY_TYPE	nnozero, 62
barray-meat-operators.hpp, 210, 212	nrow, 62
barray-meat.hpp, 216	operator*=, 63
BArrayCell	operator(), 62, 63
BArrayCell< Cell_Type, Data_Type >, 50	operator+=, 63
BArrayCell< Cell_Type, Data_Type >, 49	operator-=, 63, 64
~BArrayCell, 50	operator/=, 64
BArray< Cell_Type, Data_Type >, 48	operator=, 64
BArrayCell, 50	operator==, 64
operator Cell_Type, 50	out_of_range, 65
operator*=, 50	print, 65
operator+=, 50	reserve, 65
operator-=, 51	resize, 65
operator/=, 51	rm_cell, 65
operator=, 51	row, 65, 66
operator==, 51	rowsum, 66
BArrayCell_const	set_data, 66
BArrayCell_const< Cell_Type, Data_Type >, 52	swap_cells, 66
BArrayCell_const< Cell_Type, Data_Type >, 52	swap_cols, 67
~BArrayCell_const, 52	swap_rows, 67
BArray< Cell_Type, Data_Type >, 48	toggle_cell, 67
BArrayCell_const, 52	toggle_lock, 67
operator Cell_Type, 53	transpose, 67
operator!=, 53	visited, 69
operator<, 53	zero_col, 68
operator<=, 53	zero_row, 68
operator>, 53	barraydense-bones.hpp
operator>=, 54	BARRY_BARRAYDENSE_BONES_HPP, 235
operator==, 53	barraydense-meat-operators.hpp
BArrayDense	BARRY_BARRAYDENSE_MEAT_OPERATORS_HPP,
BArrayDense < Cell_Type, Data_Type >, 57, 58	236
BArrayDense< Cell_Type, Data_Type >, 54	BDENSE_TEMPLATE, 236–238
∼BArrayDense, 58	BDENSE_TEMPLATE_ARGS, 236, 238
BArrayDense, 57, 58	BDENSE_TYPE, 236, 238
BArrayDenseCell< Cell_Type, Data_Type >, 68,	COL, 237
72	POS, 237

POS_N, 237 ROW, 237	BArrayDenseRow< Cell_Type, Data_Type >, 79 BArrayDenseRow_const< Cell_Type, Data_Type
barraydense-meat.hpp	>, 82
add, 254	operator Cell_Type, 71
ans, 243, 254	operator*=, 71
BDENSE TEMPLATE, 241, 243–250	operator+=, 71
——————————————————————————————————————	
BDENSE_TEMPLATE_ARGS, 241	operator-=, 71
BDENSE_TYPE, 242	operator/=, 71
check_bounds, 254	operator=, 71
check_exists, 254	operator==, 72
COL, 242	barraydensecell-bones.hpp
col, 255	POS, 260
const, 255	barraydensecell-meat.hpp
copy_data, 255	POS, 262
data, 255	BArrayDenseCell_const< Cell_Type, Data_Type >, 73
delete_data, 255	BArrayDenseCol < Cell_Type, Data_Type >, 75
delete_data_, 256	BArrayDenseCol_const< Cell_Type, Data_Type >,
el, 256	77
el_colsums, 256	BArrayDenseRow< Cell_Type, Data_Type >, 79
el_rowsums, 256	BArrayDenseRow_const< Cell_Type, Data_Type
else, 256	>, 82
false, 257	BArrayDenseCol
for, 250	BArrayDenseCol< Cell_Type, Data_Type >, 73
i1, 257	BArrayDenseCol< Cell_Type, Data_Type >, 73
if, 250, 251	BArrayDense< Cell_Type, Data_Type >, 68, 75
	BArrayDenseCell< Cell_Type, Data_Type >, 00, 73
insert_cell, 251	75
j, 257	
j0, 257	BArrayDenseCell_const< Cell_Type, Data_Type
j1, 257	>, 75
M, 251, 257	BArrayDenseCol, 73
M_, 258	begin, 74
N, 258	end, 74
POS, 242	operator(), 74
POS_N, 242	size, 74
report, 258	barraydensecol-bones.hpp
resize, 252	POS, 263
return, 258	POS_N, 263
rm_cell, 253	ZERO_CELL, 263
ROW, 242	BArrayDenseCol_const
source, 258	BArrayDenseCol_const< Cell_Type, Data_Type >,
target, 259	76
v, 259	BArrayDenseCol_const< Cell_Type, Data_Type >, 75
va_end, 253	BArrayDense< Cell_Type, Data_Type >, 68
va start, 253	BArrayDenseCell< Cell_Type, Data_Type >, 72,
val0, 259	77
val1, 259	BArrayDenseCell const< Cell Type, Data Type
value, 259	>, 77
vprintf, 253	BArrayDenseCol_const, 76
ZERO_CELL, 243	begin, 76
BArrayDenseCell	end, 76
BArrayDenseCell < Cell_Type, Data_Type >, 70	operator(), 76
BArrayDenseCell Cell_Type, Data_Type >, 69	size, 77
~BArrayDenseCell, 70	BArrayDenseRow
BArrayDense< Cell_Type, Data_Type >, 68, 72	BArrayDenseRow< Cell_Type, Data_Type >, 78
BArrayDenseCell, 70	BArrayDenseRow< Cell_Type, Data_Type >, 77
BArrayDenseCol< Cell_Type, Data_Type >, 72, 75	BArrayDense < Cell_Type, Data_Type >, 69, 79
BArrayDenseCol_const< Cell_Type, Data_Type >,	BArrayDenseCell< Cell_Type, Data_Type >, 79
72, 77	

BArrayDenseCell_const< Cell_Type, Data_Type	BArrayVector, 87
>, 79	begin, 88
BArrayDenseRow, 78	end, 88
begin, 78	is_col, 88
end, 78	is_row, 88
operator(), 79	operator std::vector< Cell_Type >, 89
size, 79	operator*=, 89
barraydenserow-bones.hpp	operator+=, 89
POS, 264	operator-=, 89
POS_N, 264	operator/=, 89
ZERO_CELL, 264 BArrayDenseRow_const	operator=, 90
BArrayDenseRow_const< Cell_Type, Data_Type	operator==, 90 size, 90
>, 80	barrayvector-meat.hpp
BArrayDenseRow_const< Cell_Type, Data_Type >, 80	BARRY_BARRAYVECTOR_MEAT_HPP, 270
BArrayDense< Cell_Type, Data_Type >, 69	BArrayVector_const
BArrayDenseCell< Cell_Type, Data_Type >, 82	BArrayVector_const< Cell_Type, Data_Type >, 91
BArrayDenseCell_const< Cell_Type, Data_Type	BArrayVector_const< Cell_Type, Data_Type >, 90
>, 82	~BArrayVector const, 91
BArrayDenseRow const, 80	BArrayVector_const, 91
begin, 81	begin, 92
end, 81	end, 92
operator(), 81	is_col, 92
size, 81	is_row, 92
BArrayRow	operator std::vector< Cell_Type >, 92
BArrayRow< Cell_Type, Data_Type >, 83	operator!=, 92
BArrayRow< Cell_Type, Data_Type >, 82	operator<, 93
∼BArrayRow, 83	operator<=, 93
BArrayRow, 83	operator>, 93
operator BArrayRow< Cell_Type, Data_Type >, 83	operator>=, 93
operator*=, 83	operator==, 93
operator+=, 83	size, 94
operator-=, 83	barry, 31
operator/=, 84 operator=, 84	barry-configuration.hpp BARRY_CHECK_SUPPORT, 271
operator==, 84	BARRY_ISFINITE, 271
barrayrow-meat.hpp	BARRY_MAX_NUM_ELEMENTS, 271
BARRY_BARRAYROW_MEAT_HPP, 266	BARRY_SAFE_EXP, 271
BROW TEMPLATE, 266–268	Map, 271
BROW_TEMPLATE_ARGS, 267	printf_barry, 271
BROW TYPE, 267	barry-debug.hpp
BArrayRow_const	BARRY_DEBUG_LEVEL, 272
BArrayRow_const< Cell_Type, Data_Type >, 85	barry-macros.hpp
BArrayRow_const< Cell_Type, Data_Type >, 84	BARRY_ONE, 273
\sim BArrayRow_const, 85	BARRY_ONE_DENSE, 273
BArrayRow_const, 85	BARRY_UNUSED, 273
operator BArrayRow_const< Cell_Type, Data_Type	BARRY_ZERO, 273
>, 85	BARRY_ZERO_DENSE, 273
operator!=, 85	barry.hpp
operator<, 85	BARRY_HPP, 275
operator<=, 86	BARRY_VERSION, 275
operator>, 86	COUNTER_FUNCTION, 275
operator = , 86	COUNTER_LAMBDA, 276
operator==, 86 BArrayVector	RULE_FUNCTION, 276 RULE_LAMBDA, 276
BArray Vector < Cell_Type, Data_Type >, 87	barry::counters, 31
BArrayVector< Cell_Type, Data_Type >, 86	barry::counters; 31 barry::counters::defm, 32
~BArrayVector. 88	barry::counters::network, 32

barry::counters::phylo, 32 BARRY_BARRAY_MEAT_OPERATORS_HPP	BArrayDenseRow_const< Cell_Type, Data_Type >, 81
barray-meat-operators.hpp, 210	BArrayVector $<$ Cell_Type, Data_Type $>$, 88
BARRY_BARRAYDENSE_BONES_HPP	BArrayVector_const< Cell_Type, Data_Type >, 92
barraydense-bones.hpp, 235	PhyloCounterData, 173
BARRY_BARRAYDENSE_MEAT_OPERATORS_HPP	PowerSet < Array_Type, Data_Rule_Type >, 180
barraydense-meat-operators.hpp, 236	blengths
BARRY_BARRAYROW_MEAT_HPP	NodeData, 172
barrayrow-meat.hpp, 266	BOTH
BARRY_BARRAYVECTOR_MEAT_HPP	CHECK, 32
barrayvector-meat.hpp, 270	EXISTS, 34
BARRY_CHECK_SUPPORT	BROW_TEMPLATE
barry-configuration.hpp, 271	barrayrow-meat.hpp, 266–268
BARRY_DEBUG_LEVEL	BROW_TEMPLATE_ARGS
barry-debug.hpp, 272	barrayrow-meat.hpp, 267
BARRY HPP	BROW_TYPE
barry.hpp, 275	barrayrow-meat.hpp, 267
BARRY_ISFINITE	•
barry-configuration.hpp, 271	calc
BARRY MAX NUM ELEMENTS	PowerSet < Array_Type, Data_Rule_Type >, 180
barry-configuration.hpp, 271	Support< Array_Type, Data_Counter_Type,
BARRY_ONE	Data_Rule_Type, Data_Rule_Dyn_Type >,
barry-macros.hpp, 273	198
BARRY ONE DENSE	calc_backend_dense
	support-meat.hpp, 347
barry-macros.hpp, 273	calc_backend_sparse
BARRY_PROGRESS_BAR_WIDTH	support-meat.hpp, 348
progress.hpp, 334	calc_reduced_sequence
BARRY_SAFE_EXP	Geese, 133
barry-configuration.hpp, 271	calc_sequence
BARRY_SUPPORT_MEAT_HPP	
support-meat.hpp, 347	Geese, 133, 134
BARRY_UNUSED	Cell
barry-macros.hpp, 273	Cell< Cell_Type >, 95, 96
BARRY_VERSION	Cell< Cell_Type >, 94
barry.hpp, 275	∼Cell, 95
BARRY_ZERO	active, 98
barry-macros.hpp, 273	add, 96, 97
BARRY_ZERO_DENSE	Cell, 95, 96
barry-macros.hpp, 273	operator Cell_Type, 97
BARRY_ZERO_NETWORK	operator!=, 97
network.hpp, 304	operator=, 97, 98
BARRY_ZERO_NETWORK_DENSE	operator==, 98
network.hpp, 305	value, 98
BDENSE_TEMPLATE	visited, 98
barraydense-meat-operators.hpp, 236–238	Cell_const< Cell_Type >, 99
barraydense-meat.hpp, 241, 243–250	change_stats
BDENSE_TEMPLATE_ARGS	Support< Array_Type, Data_Counter_Type,
barraydense-meat-operators.hpp, 236, 238	Data_Rule_Type, Data_Rule_Dyn_Type >,
barraydense-meat.hpp, 241	201
BDENSE_TYPE	change_stats_different
barraydense-meat-operators.hpp, 236, 238	support-meat.hpp, 352
barraydense-meat.hpp, 242	CHECK, 32
	BOTH, 32
BArrayDonsoCol Coll Type Data Type > 74	NONE, 32
BArrayDenseCol const < Coll Type, Data_Type >, 74	ONE, 32
BArrayDenseCol_const< Cell_Type, Data_Type >,	TWO, 33
76 PArroy Dance Poys < Call Type Data Type > 70	check_bounds
BArrayDenseRow< Cell_Type, Data_Type >, 78	barray-meat.hpp, 226
	barraydense-meat.hpp, 254

check_exists	coordinates_free
barray-meat.hpp, 227	PowerSet< Array_Type, Data_Rule_Type >, 182
barraydense-meat.hpp, 254	Support< Array_Type, Data_Counter_Type,
clear	Data_Rule_Type, Data_Rule_Dyn_Type >,
BArray< Cell_Type, Data_Type >, 39	202
BArrayDense < Cell_Type, Data_Type >, 58	coordinates_locked
FreqTable < T >, 126	PowerSet < Array_Type, Data_Rule_Type >, 182
statscounter-meat.hpp, 340	Support< Array_Type, Data_Counter_Type,
COL	Data_Rule_Type, Data_Rule_Dyn_Type >,
barray-meat-operators.hpp, 211	202
barray-meat.hpp, 217, 221	copy_data
barraydense-meat-operators.hpp, 237	barray-meat.hpp, 227
barraydense-meat.hpp, 242	barraydense-meat.hpp, 255
col	count
BArray< Cell_Type, Data_Type >, 39	Counter< Array_Type, Data_Type >, 104
BArrayDense< Cell_Type, Data_Type >, 58, 59	count_all
barraydense-meat.hpp, 255	StatsCounter< Array_Type, Data_Type >, 192
col0	count_current
barray-meat.hpp, 227	StatsCounter< Array_Type, Data_Type >, 193
Col_type	count_fun
typedefs.hpp, 357	Counter< Array_Type, Data_Type >, 105
colnames	counters-meat.hpp, 283
Flock, 120	count_fun_
Geese, 134	counters-meat.hpp, 287
Model < Array_Type, Data_Counter_Type,	count_init
Data_Rule_Type, Data_Rule_Dyn_Type >,	StatsCounter< Array_Type, Data_Type >, 193
153	Counter
colsum	Counter< Array_Type, Data_Type >, 102, 103
BArrayDense< Cell_Type, Data_Type >, 59	counter
conditional_prob	counters-meat.hpp, 287
Model< Array_Type, Data_Counter_Type,	statscounter-meat.hpp, 342
Data_Rule_Type, Data_Rule_Dyn_Type >,	Counter< Array_Type, Data_Type >, 101
153	~Counter, 103
const	count, 104
barray-meat.hpp, 227	count_fun, 105
barraydense-meat.hpp, 255	Counter, 102, 103
ConstBArrayRowlter	data, 105
ConstBArrayRowlter< Cell_Type, Data_Type >,	desc, 105
100	get_description, 104
ConstBArrayRowlter< Cell_Type, Data_Type >, 99	get_name, 104
~ConstBArrayRowIter, 100	init, 104
Array, 100	init fun, 105
ConstBArrayRowlter, 100	name, 105
current_col, 101	operator=, 104
current_row, 101	counter
iter, 101	counters-meat.hpp, 287
coord_i	counter_absdiff
support-meat.hpp, 353	DEFMArray counters, 14
coord_i	counter_co_opt
support-meat.hpp, 353	Phylo counters, 23
coordiantes_n_free	counter_cogain
Support< Array_Type, Data_Counter_Type,	Phylo counters, 23
Data_Rule_Type, Data_Rule_Dyn_Type >,	counter_css_census01
202	network-css.hpp, 297
coordiantes_n_locked	counter_css_census02
Support< Array_Type, Data_Counter_Type,	network-css.hpp, 298
Data_Rule_Type, Data_Rule_Dyn_Type >,	counter_css_census03
202	network-css.hpp, 298

counter_css_census04	counter_k_genes_changing
network-css.hpp, 298	Phylo counters, 25
counter_css_census05	COUNTER_LAMBDA
network-css.hpp, 298	barry.hpp, 276
counter_css_census06	counter_less_than_p_prop_genes_changing
network-css.hpp, 299	Phylo counters, 25
counter_css_census07	counter_longest
network-css.hpp, 299	Phylo counters, 25
counter_css_census08	counter_loss
network-css.hpp, 299	Phylo counters, 25
counter_css_census09	counter maxfuns
network-css.hpp, 299	Phylo counters, 26
counter_css_census10	counter_mutual
network-css.hpp, 300	DEFMArray counters, 17
• •	
counter_css_completely_false_recip_comiss	counter_neofun
network-css.hpp, 300	Phylo counters, 26
counter_css_completely_false_recip_omiss	counter_neofun_a2b
network-css.hpp, 300	Phylo counters, 26
counter_css_mixed_recip	counter_nodecov
network-css.hpp, 300	DEFMArray counters, 18
counter_css_partially_false_recip_commi	counter_nodeicov
network-css.hpp, 301	DEFMArray counters, 18
counter_css_partially_false_recip_omiss	counter_nodematch
network-css.hpp, 301	DEFMArray counters, 18
counter_ctriads	counter_nodeocov
DEFMArray counters, 14	DEFMArray counters, 18
counter_degree	counter_odegree
DEFMArray counters, 14	DEFMArray counters, 18, 19
counter_deleted	counter_odegree15
statscounter-meat.hpp, 342	DEFMArray counters, 19
counter_density	counter_ones
DEFMArray counters, 15	DEFMArray counters, 19
counter diff	counter_ostar2
DEFMArray counters, 15	DEFMArray counters, 20
counter edges	counter overall changes
DEFMArray counters, 15	Phylo counters, 26
counter_fixed_effect	counter_overall_gains
DEFMArray counters, 15	Phylo counters, 27
Counter_fun_type	counter_overall_gains_from_0
typedefs.hpp, 358	Phylo counters, 27
COUNTER FUNCTION	counter_overall_loss
barry.hpp, 275	Phylo counters, 27
counter gains	
	counter_pairwise_first_gain Phylo counters, 27
Phylo counters, 24	
counter_gains_from_0	counter_pairwise_neofun_singlefun
Phylo counters, 24	Phylo counters, 28
counter_gains_k_offspring	counter_pairwise_overall_change
Phylo counters, 24	Phylo counters, 28
counter_genes_changing	counter_pairwise_preserving
Phylo counters, 24	Phylo counters, 28
counter_idegree	counter_preserve_pseudogene
DEFMArray counters, 16	Phylo counters, 28
counter_idegree15	counter_prop_genes_changing
DEFMArray counters, 16	Phylo counters, 29
counter_isolates	counter_subfun
DEFMArray counters, 17	Phylo counters, 29
counter_istar2	COUNTER_TEMPLATE
DEFMArray counters, 17	counters-meat.hpp, 281, 283, 284

COUNTER_TEMPLATE_ARGS	counts
counters-meat.hpp, 282	PhyloRuleDynData, 176
counter_transition	Counts_type
DEFMArray counters, 20	typedefs.hpp, 358
counter_ttriads	covariates
DEFMArray counters, 21	DEFMData, 114
COUNTER_TYPE	CSS_APPEND
counters-meat.hpp, 282	network-css.hpp, 295
Counters	CSS_CASE_ELSE
Counters < Array_Type, Data_Type >, 107	network-css.hpp, 295
counters-meat.hpp, 284	CSS_CASE_PERCEIVED
counters	network-css.hpp, 296
statscounter-meat.hpp, 342	CSS_CASE_TRUTH
support-meat.hpp, 353	network-css.hpp, 296
Counters< Array_Type, Data_Type >, 106	CSS_CHECK_SIZE
~Counters, 107	network-css.hpp, 296
add counter, 108	CSS CHECK SIZE INIT
Counters, 107	network-css.hpp, 296
get_descriptions, 108	CSS_NET_COUNTER_LAMBDA_INIT
get_names, 108	network-css.hpp, 296
operator=, 108, 109	CSS_PERCEIVED_CELLS
operator[], 109	network-css.hpp, 297
size, 109	CSS_SIZE
counters-meat.hpp	network-css.hpp, 297
count_fun, 283	CSS_TRUE_CELLS
count_fun_, 287	network-css.hpp, 297
counter, 287	current_col
counter_, 287	ConstBArrayRowlter< Cell_Type, Data_Type >,
COUNTER_TEMPLATE, 281, 283, 284	101
COUNTER_TEMPLATE_ARGS, 282	current_row
COUNTER_TYPE, 282	ConstBArrayRowlter< Cell_Type, Data_Type >,
Counters, 284	101
COUNTERS_TEMPLATE, 282, 284, 285	current_stats
COUNTERS_TEMPLATE_ARGS, 282	statscounter-meat.hpp, 342
COUNTERS_TYPE, 282	Support< Array_Type, Data_Counter_Type,
data, 286	Data_Rule_Type, Data_Rule_Dyn_Type >,
data_, 287	202
desc, 286	D
desc_, 288	BArray< Cell Type, Data Type >, 40
i, 288	BArrayDense< Cell Type, Data Type >, 59
init_fun, 286	
init_fun_, 288	Rule < Array_Type, Data_Type >, 187
j, 288	D_ptr
name, 286	BArray Cell_Type, Data_Type >, 40
name_, 288	BArrayDense < Cell_Type, Data_Type >, 59
noexcept, 289	dat
return, 289	Flock, 123
this, 289	data
counters_	barray-meat.hpp, 228
statscounter-meat.hpp, 342	barraydense-meat.hpp, 255
support-meat.hpp, 353	Counter< Array_Type, Data_Type >, 105
COUNTERS_TEMPLATE	counters-meat.hpp, 286
counters-meat.hpp, 282, 284, 285	PowerSet < Array_Type, Data_Rule_Type >, 182
COUNTERS_TEMPLATE_ARGS	data_
counters-meat.hpp, 282	counters-meat.hpp, 287
COUNTERS_TYPE	DEFAULT_DUPLICATION
counters-meat.hpp, 282	phylo.hpp, 311
Counting, 11	default_val
.	BArray< Cell_Type, Data_Type >, 40

BArrayDense < Cell_Type, Data_Type >, 60	DEFMCounterData, 110, 111
defm-bones.hpp	idx, 111
INITIALIZED, 322	indices, 111
keygen_full, 322	markov_order, 112
RULE_FUNCTION, 322	num, 111
vec_diff, 322	numbers, 112
defm.hpp	DEFMCounters
DEFM COUNTER, 291	defm.hpp, 292
DEFM_COUNTER_LAMBDA, 291	DEFMData, 112
DEFM RULE, 291	∼DEFMData, 114
DEFM RULE LAMBDA, 291	at, 114
DEFMArray, 292	covariates, 114
DEFMCounter, 292	DEFMData, 113
DEFMCounters, 292	n_covariates, 115
DEFMModel, 292	obs_n_times, 115
DEFMRule, 292	obs_start, 115
DEFMRules, 293	operator(), 114
DEFMStatsCounter, 293	DEFMModel
DEFMSupport, 293	defm.hpp, 292
DEFM COUNTER	DEFMRule
defm.hpp, 291	defm.hpp, 292
DEFM_COUNTER_LAMBDA	DEFMRules
defm.hpp, 291	defm.hpp, 293
DEFM_RULE	DEFMStatsCounter
defm.hpp, 291	defm.hpp, 293
DEFM_RULE_LAMBDA	DEFMSupport
	• •
defm.hpp, 291	defm.hpp, 293
DEFMArray	delete_counters
defm.hpp, 292	Support< Array_Type, Data_Counter_Type,
DEFMArray counters, 12	Data_Rule_Type, Data_Rule_Dyn_Type >,
counter_absdiff, 14	203
counter_ctriads, 14	support-meat.hpp, 353
counter_degree, 14	delete_data
counter_density, 15	barray-meat.hpp, 228
counter_diff, 15	barraydense-meat.hpp, 255
counter_edges, 15	delete_data_
counter_fixed_effect, 15	barray-meat.hpp, 228
counter_idegree, 16	barraydense-meat.hpp, 256
counter_idegree15, 16	delete_rengine
counter_isolates, 17	Geese, 145
counter_istar2, 17	delete_rules
counter_mutual, 17	Support< Array_Type, Data_Counter_Type,
counter_nodecov, 18	Data_Rule_Type, Data_Rule_Dyn_Type >,
counter_nodeicov, 18	203
counter_nodematch, 18	support-meat.hpp, 353
counter_nodeocov, 18	delete_rules_dyn
counter_odegree, 18, 19	Support< Array_Type, Data_Counter_Type,
counter_odegree15, 19	
counter_ones, 19	Data_Rule_Type, Data_Rule_Dyn_Type >,
	Data_Rule_Type, Data_Rule_Dyn_Type >, 203
counter_ostar2, <mark>20</mark>	
counter_ostar2, 20 counter_transition, 20	203
	203 support-meat.hpp, 354
counter_transition, 20	203 support-meat.hpp, 354 delete_support
counter_transition, 20 counter_ttriads, 21 NETWORK_COUNTER, 21	203 support-meat.hpp, 354 delete_support Geese, 146 desc
counter_transition, 20 counter_triads, 21	203 support-meat.hpp, 354 delete_support Geese, 146 desc Counter< Array_Type, Data_Type >, 105
counter_transition, 20 counter_ttriads, 21 NETWORK_COUNTER, 21 rules_zerodiag, 21 DEFMCounter	203 support-meat.hpp, 354 delete_support Geese, 146 desc Counter< Array_Type, Data_Type >, 105 counters-meat.hpp, 286
counter_transition, 20 counter_ttriads, 21 NETWORK_COUNTER, 21 rules_zerodiag, 21	203 support-meat.hpp, 354 delete_support Geese, 146 desc Counter< Array_Type, Data_Type >, 105

NetworkData, 165	UKNOWN, 34
DUPL DUPL	
phylo.hpp, 311	f_
DUPL EITH	statscounter-meat.hpp, 343
phylo.hpp, 311	support-meat.hpp, 354
DUPL_SPEC	false
phylo.hpp, 311	barray-meat.hpp, 228
duplication	barraydense-meat.hpp, 257
Node, 169	first
NodeData, 172	barray-meat.hpp, 229
PhyloRuleDynData, 176	Flock, 118
	\sim Flock, 119
el	add_data, 119
barraydense-meat.hpp, 256	colnames, 120
el_colsums	dat, 123
barraydense-meat.hpp, 256	Flock, 119
el_rowsums	get_counters, 120
barraydense-meat.hpp, 256	get_model, 120
else	get_stats_support, 120
barray-meat.hpp, 228	get_stats_target, 120
barraydense-meat.hpp, 256	get_support_fun, 120
support-meat.hpp, 354	init, 121
empty	initialized, 124
PhyloCounterData, 173	likelihood_joint, 121
EmptyArray	model, 124
PowerSet < Array_Type, Data_Rule_Type >, 182	nfunctions, 124
statscounter-meat.hpp, 343	nfuns, 121
end	nleafs, 121
BArrayDenseCol Cell_Type, Data_Type >, 74	nnodes, 122
BArrayDenseCol_const< Cell_Type, Data_Type >,	nterms, 122
76	ntrees, 122
BArrayDenseRow< Cell_Type, Data_Type >, 78	operator(), 122
BArrayDenseRow_const< Cell_Type, Data_Type	parse_polytomies, 123
>, 81	print, 123
BArrayVector< Cell_Type, Data_Type >, 88 BArrayVector_const< Cell_Type, Data_Type >, 92	rengine, 124
PhyloCounterData, 174	set_seed, 123 support_size, 123
PowerSet < Array_Type, Data_Rule_Type >, 180	flush_data
Progress, 185	BArray < Cell_Type, Data_Type >, 40
Entries	for
Entries < Cell_Type >, 116	barray-meat-operators.hpp, 212
Entries < Cell_Type >, 115	barray-meat.hpp, 222
~Entries, 116	barraydense-meat.hpp, 250
Entries, 116	statscounter-meat.hpp, 340
resize, 117	support-meat.hpp, 348
source, 117	FreqTable
target, 117	FreqTable $<$ T $>$, 125
val, 117	FreqTable < T >, 124
eval_rules_dyn	\sim FreqTable, 125
Support< Array_Type, Data_Counter_Type,	add, 126
Data_Rule_Type, Data_Rule_Dyn_Type >,	as_vector, 126
199	clear, 126
EXISTS, 33	FreqTable, 125
AS_ONE, 33	get_data, 126
AS_ZERO, 33	get_index, 126
BOTH, 34	make_hash, 127
NONE, 34	print, 127
ONE, 34	reserve, 127
TWO, 34	size, 127

Geese, 128	get_arrays2support
∼Geese, 132, 133	Model < Array_Type, Data_Counter_Type.
calc_reduced_sequence, 133	Data_Rule_Type, Data_Rule_Dyn_Type >,
calc_sequence, 133, 134	154
colnames, 134	get cell
delete_rengine, 145	BArray< Cell_Type, Data_Type >, 40
delete_support, 146	BArrayDense < Cell_Type, Data_Type >, 60
Geese, 131–133	get_col_vec
get annotated nodes, 134	BArray< Cell_Type, Data_Type >, 41
get_counters, 134, 135	BArrayDense < Cell_Type, Data_Type >, 60
get model, 135	get counters
get_probabilities, 135	Flock, 120
get_rengine, 135	Geese, 134, 135
get_states, 136	Model < Array_Type, Data_Counter_Type,
get_support_fun, 136	Data_Rule_Type, Data_Rule_Dyn_Type >:
	154
inherit_support, 136, 137	
init, 137	PhyloCounterData, 174
init_node, 137	StatsCounter< Array_Type, Data_Type >, 193
initialized, 146	Support< Array_Type, Data_Counter_Type
likelihood, 137, 138	Data_Rule_Type, Data_Rule_Dyn_Type >
likelihood_exhaust, 138	199
map_to_nodes, 146	get_counts
nannotations, 138	Support< Array_Type, Data_Counter_Type,
nfunctions, 146	Data_Rule_Type, Data_Rule_Dyn_Type >
nfuns, 139	199
nleafs, 139	get_current_stats
nnodes, 139	Support< Array_Type, Data_Counter_Type
nodes, 146	Data_Rule_Type, Data_Rule_Dyn_Type >
nterms, 140	199
observed_counts, 140	get_data
operator=, 140, 141	BArrayDense< Cell_Type, Data_Type >, 60
parse_polytomies, 141	FreqTable $< T >$, 126
predict, 141	PowerSet< Array_Type, Data_Rule_Type >, 180
predict_backend, 142	Support< Array_Type, Data_Rounter_Type,
predict_exhaust, 142	Data_Rule_Type, Data_Rule_Dyn_Type >:
•	199
predict_exhaust_backend, 142, 143	
predict_sim, 143	get_data_ptr
print, 143	PowerSet< Array_Type, Data_Rule_Type >, 181
print_observed_counts, 144	get_description
pset_loc, 146	Counter< Array_Type, Data_Type >, 104
reduced_sequence, 147	get_descriptions
sequence, 147	Counters < Array_Type, Data_Type >, 108
set_seed, 144	StatsCounter< Array_Type, Data_Type >, 193
simulate, 144	get_entries
support_size, 145	BArray< Cell_Type, Data_Type >, 41
update_annotations, 145	BArrayDense < Cell_Type, Data_Type >, 60
geese-bones.hpp	get_index
INITIALIZED, 325	FreqTable $<$ T $>$, 126
keygen_full, 326	get_last_name
RULE_FUNCTION, 326	phylo.hpp, 316
vec_diff, 326	get_model
vector_caster, 326	Flock, 120
gen_key	Geese, 135
Model< Array_Type, Data_Counter_Type,	get_name
Data_Rule_Type, Data_Rule_Dyn_Type >,	Counter< Array_Type, Data_Type >, 104
153	get_names
get_annotated_nodes	Counters < Array_Type, Data_Type >, 108
Geese, 134	StatsCounter< Array_Type, Data_Type >, 100
GEESE, IUT	olalooounter < Array_Type, Dala_Type >, 193

get norm const	Data_Rule_Type, Data_Rule_Dyn_Type >
Model Array_Type, Data_Counter_Type,	156
2= 31 / = = 31 /	
Data_Rule_Type, Data_Rule_Dyn_Type >,	
154	Flock, 120
get_parent	Geese, 136
Node, 168	Model< Array_Type, Data_Counter_Type
get_probabilities	Data_Rule_Type, Data_Rule_Dyn_Type >
Geese, 135	156
get_pset	
Model < Array_Type, Data_Counter_Type,	hashes
Data_Rule_Type, Data_Rule_Dyn_Type >,	Support< Array_Type, Data_Counter_Type
154	$Data_Rule_Type, Data_Rule_Dyn_Type >$
get_pset_arrays	203
Model< Array_Type, Data_Counter_Type,	support-meat.hpp, 354
Data_Rule_Type, Data_Rule_Dyn_Type >,	hashes_initialized
154	Support< Array_Type, Data_Counter_Type
get_pset_probs	Data_Rule_Type, Data_Rule_Dyn_Type >
Model < Array_Type, Data_Counter_Type,	203
Data_Rule_Type, Data_Rule_Dyn_Type >,	
	i
155	counters-meat.hpp, 288
get_pset_stats	i1
Model< Array_Type, Data_Counter_Type,	barray-meat.hpp, 229
Data_Rule_Type, Data_Rule_Dyn_Type >,	barraydense-meat.hpp, 257
155	id
get_rengine	
Geese, 135	Node, 169
Model< Array_Type, Data_Counter_Type,	idx
Data_Rule_Type, Data_Rule_Dyn_Type >,	DEFMCounterData, 111
155	if
get_row_vec	barray-meat.hpp, 222–225
BArray< Cell_Type, Data_Type >, 41	barraydense-meat.hpp, 250, 251
BArrayDense< Cell_Type, Data_Type >, 61	support-meat.hpp, 348
get_rules	IF_MATCHES
Model< Array_Type, Data_Counter_Type,	phylo.hpp, 312
Data_Rule_Type, Data_Rule_Dyn_Type >,	IF_NOTMATCHES
155	phylo.hpp, 312
Support< Array_Type, Data_Counter_Type,	include/barry/barray-bones.hpp, 207
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/barray-iterator.hpp, 208
200	include/barry/barray-meat-operators.hpp, 209
	include/barry/barray-meat.hpp, 214
get_rules_dyn	include/barry/barraycell-bones.hpp, 232
Model < Array_Type, Data_Counter_Type,	include/barry/barraycell-meat.hpp, 233
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/barraydense-bones.hpp, 233
156	include/barry/barraydense-meat-operators.hpp, 235
Support< Array_Type, Data_Counter_Type,	include/barry/barraydense-meat.hpp, 239
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/barraydensecell-bones.hpp, 260
200	
get_seq	include/barry/barraydensecell-meat.hpp, 261
Rules< Array_Type, Data_Type >, 189	include/barry/barraydensecol-bones.hpp, 262
get_states	include/barry/barraydenserow-bones.hpp, 263
Geese, 136	include/barry/barrayrow-bones.hpp, 265
get_stats_support	include/barry/barrayrow-meat.hpp, 265
Flock, 120	include/barry/barrayvector-bones.hpp, 268
Model< Array_Type, Data_Counter_Type,	include/barry/barrayvector-meat.hpp, 269
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/barry-configuration.hpp, 270
156	include/barry/barry-debug.hpp, 272
get_stats_target	include/barry/barry-macros.hpp, 272
Flock, 120	include/barry/barry.hpp, 274
	include/barry/cell-bones.hpp, 277
Model< Array_Type, Data_Counter_Type,	include/barry/cell-meat.hpp, 277

include/barry/col-bones.hpp, 278	init_node
include/barry/counters-bones.hpp, 278	Geese, 137
include/barry/counters-meat.hpp, 280	init_support
include/barry/counters/defm.hpp, 289	PowerSet < Array_Type, Data_Rule_Type >, 181
include/barry/counters/network-css.hpp, 294	Support< Array_Type, Data_Counter_Type,
include/barry/counters/network.hpp, 302	Data_Rule_Type, Data_Rule_Dyn_Type >,
include/barry/counters/phylo.hpp, 309	200
include/barry/freqtable.hpp, 316	INITIALIZED
include/barry/model-bones.hpp, 317	defm-bones.hpp, 322
include/barry/model-meat.hpp, 319	geese-bones.hpp, 325
include/barry/models/defm.hpp, 293	initialized
include/barry/models/defm/defm-bones.hpp, 321	Flock, 124
include/barry/models/defm/defm-meat.hpp, 323	Geese, 146
include/barry/models/geese.hpp, 323	insert_cell
include/barry/models/geese/flock-bones.hpp, 324	BArray< Cell_Type, Data_Type >, 42
include/barry/models/geese/flock-meat.hpp, 324	BArrayDense< Cell_Type, Data_Type >, 61
include/barry/models/geese/geese-bones.hpp, 325	barraydense-meat.hpp, 251
include/barry/models/geese/geese-meat-constructors.hpp	support-meat.hpp, 348, 349
327	is_col
include/barry/models/geese/geese-meat-likelihood.hpp,	BArrayVector< Cell_Type, Data_Type >, 88
327	BArrayVector_const< Cell_Type, Data_Type >, 92
include/barry/models/geese/geese-meat-likelihood_exhau	s <u>th</u> ppnse
328	BArray< Cell_Type, Data_Type >, 42
include/barry/models/geese/geese-meat-predict.hpp,	BArrayDense< Cell_Type, Data_Type >, 62
329	IS_DUPLICATION
include/barry/models/geese/geese-meat-predict_exhaust.l	hpp, phylo.hpp, 312
329	IS_EITHER
include/barry/models/geese/geese-meat-predict_sim.hpp,	phylo.hpp, 312
330	is_empty
include/barry/models/geese/geese-meat-simulate.hpp,	BArray< Cell_Type, Data_Type >, 42
330	BArrayDense< Cell_Type, Data_Type >, 62
include/barry/models/geese/geese-meat.hpp, 331	is_leaf
include/barry/models/geese/geese-node-bones.hpp,	Node, 168
331	is_row
include/barry/powerset-bones.hpp, 332	BArrayVector< Cell_Type, Data_Type >, 88
include/barry/powerset-meat.hpp, 333	BArrayVector_const< Cell_Type, Data_Type >, 92
include/barry/progress.hpp, 334	IS_SPECIATION
include/barry/rules-bones.hpp, 334	phylo.hpp, 312
include/barry/rules-meat.hpp, 336	iter
include/barry/statscounter-bones.hpp, 336	ConstBArrayRowIter< Cell_Type, Data_Type >,
include/barry/statscounter-meat.hpp, 338	101
include/barry/support-bones.hpp, 344	
include/barry/support-meat.hpp, 345	j
include/barry/typedefs.hpp, 356	barray-meat.hpp, 229
indices	barraydense-meat.hpp, 257
DEFMCounterData, 111	counters-meat.hpp, 288
NetCounterData, 163	statscounter-meat.hpp, 343
inherit_support	j0
Geese, 136, 137	barray-meat.hpp, 229
init	barraydense-meat.hpp, 257
7= 71 / = 71 /	j1
Flock, 121	barray-meat.hpp, 229
Geese, 137	barraydense-meat.hpp, 257
init_fun	keygen_default
Counter< Array_Type, Data_Type >, 105	model-bones.hpp, 319
counters-meat.hpp, 286	
init_fun_	keygen_full defm-bones.hpp, 322
counters-meat.hpp, 288	·
	geese-bones.hpp, 326

lb	gen_key, 153
PhyloRuleDynData, 176	get_arrays2support, 154
likelihood	get_counters, 154
Geese, 137, 138	get_norm_const, 154
Model< Array_Type, Data_Counter_Type,	get_pset, 154
Data_Rule_Type, Data_Rule_Dyn_Type >,	get_pset_arrays, 154
156, 157	get_pset_probs, 155
likelihood_	get_pset_stats, 155
model-meat.hpp, 320	get_rengine, 155
likelihood_exhaust	get_rules, 155
Geese, 138	get_rules_dyn, 156
likelihood_joint	get_stats_support, 156
Flock, 121	get_stats_target, 156
likelihood_total	get_support_fun, 156
Model< Array_Type, Data_Counter_Type,	likelihood, 156, 157
Data_Rule_Type, Data_Rule_Dyn_Type >,	likelihood_total, 157
157	Model, 150, 151
	nterms, 158
M	operator=, 158
barray-meat.hpp, 225, 229	print, 158
barraydense-meat.hpp, 251, 257	print_stats, 158
PowerSet < Array_Type, Data_Rule_Type >, 182	sample, 158, 159
Support Array_Type, Data_Counter_Type,	set_counters, 159
Data_Rule_Type, Data_Rule_Dyn_Type >,	set_keygen, 159
204	set_rengine, 159
M_	set_rules, 159
barray-meat.hpp, 230	set_rules_dyn, 160
barraydense-meat.hpp, 258	set_seed, 160
MAKE_DUPL_VARS	set_transform_model, 160
phylo.hpp, 313	size, 161
make_hash	size_unique, 161
FreqTable < T >, 127	store_psets, 161
Map	support_size, 161
barry-configuration.hpp, 271	transform_model, 161
map_to_nodes	model-bones.hpp
Geese, 146	keygen_default, 319
MapVec_type	model-meat.hpp
typedefs.hpp, 358	likelihood_, 320
markov_order	MODEL_TEMPLATE, 320, 321
DEFMCounterData, 112	MODEL_TEMPLATE_ARGS, 320
max_num_elements	MODEL_TYPE, 320
Support	update_normalizing_constant, 321
Data_Rule_Type, Data_Rule_Dyn_Type >,	MODEL_TEMPLATE
204 Model	model-meat.hpp, 320, 321
	MODEL_TEMPLATE_ARGS
Model< Array_Type, Data_Counter_Type, Data Rule Type, Data Rule Dyn Type >,	model-meat.hpp, 320
150, 151	MODEL_TYPE
model	model-meat.hpp, 320
	N I
Flock, 124 Model < Array Type Data Counter Type Data Bule Type	N
Model < Array_Type, Data_Counter_Type, Data_Rule_Typ	11.7
Data_Rule_Dyn_Type >, 147	barraydense-meat.hpp, 258
~Model, 151	PowerSet < Array_Type, Data_Rule_Type >, 183
add_array, 151 add_counter, 152	Support< Array_Type, Data_Counter_Type
	Data_Rule_Type, Data_Rule_Dyn_Type >
add_rule, 152	204
add_rule_dyn, 152	n_counters
colnames, 153	
conditional_prob, 153	

Support< Array_Type, Data_Counter_Type,	counter_css_census10, 300
Data_Rule_Type, Data_Rule_Dyn_Type >,	counter_css_completely_false_recip_comiss, 300
204	counter_css_completely_false_recip_omiss, 300
n_covariates	counter_css_mixed_recip, 300
DEFMData, 115	counter_css_partially_false_recip_commi, 301
n free	counter_css_partially_false_recip_omiss, 301
PowerSet< Array_Type, Data_Rule_Type >, 183	CSS APPEND, 295
n locked	CSS_CASE_ELSE, 295
PowerSet< Array_Type, Data_Rule_Type >, 183	CSS_CASE_PERCEIVED, 296
name	CSS CASE TRUTH, 296
Counter< Array_Type, Data_Type >, 105	CSS CHECK SIZE, 296
counters-meat.hpp, 286	CSS_CHECK_SIZE_INIT, 296
name	CSS_NET_COUNTER_LAMBDA_INIT, 296
counters-meat.hpp, 288	CSS_PERCEIVED_CELLS, 297
nannotations	CSS_SIZE, 297
Geese, 138	CSS_TRUE_CELLS, 297
narray	network.hpp
Node, 169	BARRY_ZERO_NETWORK, 304
NCells	BARRY ZERO NETWORK DENSE, 305
barray-meat.hpp, 230	NET_C_DATA_IDX, 305
ncol	NET_C_DATA_NUM, 305
BArray< Cell_Type, Data_Type >, 43	NetCounter, 307
BArrayDense< Cell Type, Data Type >, 62	NetCounters, 307
NET_C_DATA_IDX	NetModel, 307
network.hpp, 305	NetRule, 307
NET_C_DATA_NUM	NetRules, 307
network.hpp, 305	NetStatsCounter, 307
NetCounter	
	NetSupport, 308
network.hpp, 307	Network, 308
NetCounterData, 162	NETWORK_COUNTER_LAMPRA_205
~NetCounterData, 162	NETWORK_COUNTER_LAMBDA, 305
indices, 163	NETWORK_RULE, 306
NetCounterData, 162	NETWORK_RULE_LAMBDA, 306
numbers, 163	NetworkDense, 308
NetCounters	NETWORKDENSE_COUNTER_LAMBDA, 306
network.hpp, 307	rules_zerodiag, 308
NetModel	NETWORK_COUNTER
network.hpp, 307	DEFMArray counters, 21
NetRule	network.hpp, 305
network.hpp, 307	NETWORK_COUNTER_LAMBDA
NetRules	network.hpp, 305
network.hpp, 307	NETWORK_RULE
NetStatsCounter	network.hpp, 306
network.hpp, 307	NETWORK_RULE_LAMBDA
NetSupport	network.hpp, 306
network.hpp, 308	NetworkData, 163
Network	∼NetworkData, 165
network.hpp, 308	directed, 165
network-css.hpp	NetworkData, 164
counter_css_census01, 297	vertex_attr, 165
counter_css_census02, 298	NetworkDense
counter_css_census03, 298	network.hpp, 308
counter_css_census04, 298	NETWORKDENSE_COUNTER_LAMBDA
counter_css_census05, 298	network.hpp, 306
counter_css_census06, 299	next
counter_css_census07, 299	Progress, 185
counter_css_census08, 299	nfunctions
counter_css_census09, 299	Flock, 124

Geese, 146	numbers
nfuns	DEFMCounterData, 112
Flock, 121	NetCounterData, 163
Geese, 139	
nleafs	obs_n_times
Flock, 121	DEFMData, 115
Geese, 139	obs_start
	DEFMData, 115
nnodes	observed_counts
Flock, 122	Geese, 140
Geese, 139	offspring
nnozero	Node, 169
BArray< Cell_Type, Data_Type >, 43	ONE
BArrayDense < Cell_Type, Data_Type >, 62	
Node, 165	CHECK, 32
\sim Node, 167	EXISTS, 34
annotations, 168	operator BArrayRow< Cell_Type, Data_Type >
array, 168	BArrayRow< Cell_Type, Data_Type >, 83
arrays, 169	operator BArrayRow_const< Cell_Type, Data_Type >
duplication, 169	BArrayRow_const< Cell_Type, Data_Type >, 85
get_parent, 168	operator Cell_Type
id, 169	BArrayCell< Cell_Type, Data_Type >, 50
is_leaf, 168	BArrayCell_const< Cell_Type, Data_Type >, 53
narray, 169	BArrayDenseCell< Cell_Type, Data_Type >, 71
Node, 166, 167	Cell< Cell_Type >, 97
noffspring, 168	operator std::vector< Cell_Type >
offspring, 169	BArrayVector< Cell_Type, Data_Type >, 89
, -	BArrayVector_const< Cell_Type, Data_Type >, 92
ord, 170	operator!=
parent, 170	BArrayCell_const< Cell_Type, Data_Type >, 53
probability, 170	BArrayRow_const< Cell_Type, Data_Type >, 85
subtree_prob, 170	
visited, 170	BArrayVector_const< Cell_Type, Data_Type >, 92
NodeData, 171	Cell< Cell_Type >, 97
blengths, 172	operator<
duplication, 172	BArrayCell_const< Cell_Type, Data_Type >, 53
NodeData, 171	BArrayRow_const< Cell_Type, Data_Type >, 85
states, 172	BArrayVector_const< Cell_Type, Data_Type >, 93
nodes	operator<=
Geese, 146	BArrayCell_const< Cell_Type, Data_Type >, 53
noexcept	BArrayRow_const< Cell_Type, Data_Type >, 86
counters-meat.hpp, 289	BArrayVector_const< Cell_Type, Data_Type >, 93
noffspring	operator>
Node, 168	BArrayCell_const< Cell_Type, Data_Type >, 53
NONE	BArrayRow_const< Cell_Type, Data_Type >, 86
CHECK, 32	BArrayVector_const< Cell_Type, Data_Type >, 93
	operator>=
EXISTS, 34	BArrayCell_const< Cell_Type, Data_Type >, 54
nrow	BArrayRow_const< Cell_Type, Data_Type >, 86
BArray< Cell_Type, Data_Type >, 43	BArrayVector const< Cell Type, Data Type >, 93
BArrayDense < Cell_Type, Data_Type >, 62	
nterms	operator*=
Flock, 122	BArray Cell_Type, Data_Type >, 43
Geese, 140	BArrayCell< Cell_Type, Data_Type >, 50
Model< Array_Type, Data_Counter_Type,	BArrayDense < Cell_Type, Data_Type >, 63
Data_Rule_Type, Data_Rule_Dyn_Type >,	BArrayDenseCell< Cell_Type, Data_Type >, 71
158	BArrayRow< Cell_Type, Data_Type >, 83
ntrees	BArrayVector< Cell_Type, Data_Type >, 89
Flock, 122	operator()
num	BArray< Cell_Type, Data_Type >, 43
DEFMCounterData, 111	barray-meat-operators.hpp, 213
==:	BArrayDense < Cell_Type, Data_Type >, 62, 63

	BArrayDenseCol< Cell_Type, Data_Type >, 74 BArrayDenseCol_const< Cell_Type, Data_Type >,	Cell< Cell_Type >, 98 operator[]
	76	Counters < Array_Type, Data_Type >, 109
	BArrayDenseRow< Cell_Type, Data_Type >, 79	PhyloCounterData, 174
	BArrayDenseRow_const< Cell_Type, Data_Type	PowerSet < Array_Type, Data_Rule_Type >, 181
	>, 81	ord
	DEFMData, 114	Node, 170
	Flock, 122	out_of_range
	PhyloCounterData, 174	BArray< Cell_Type, Data_Type >, 45
	Rule < Array_Type, Data_Type >, 187	BArrayDense< Cell_Type, Data_Type >, 65
	Rules< Array_Type, Data_Type >, 190	
	vecHasher< T >, 205	parent
opei	rator+=	Node, 170
- -	BArray< Cell_Type, Data_Type >, 44	parse_polytomies
	BArrayCell< Cell_Type, Data_Type >, 50	Flock, 123
	BArrayDense< Cell_Type, Data_Type >, 63	Geese, 141
	BArrayDenseCell< Cell_Type, Data_Type >, 71	Phylo counters, 22
	BArrayRow< Cell_Type, Data_Type >, 83	counter_co_opt, 23
	BArrayVector< Cell Type, Data Type >, 89	counter_cogain, 23
		counter_gains, 24
opei	rator-=	counter_gains_from_0, 24
	BArray < Cell_Type, Data_Type >, 44	counter_gains_k_offspring, 24
	BArrayCell< Cell_Type, Data_Type >, 51	counter_genes_changing, 24
	BArrayDense < Cell_Type, Data_Type >, 63, 64	counter_k_genes_changing, 25
	BArrayDenseCell< Cell_Type, Data_Type >, 71	
	BArrayRow< Cell_Type, Data_Type >, 83	counter_less_than_p_prop_genes_changing, 25
	BArrayVector< Cell_Type, Data_Type >, 89	counter_longest, 25
ope	rator/=	counter_loss, 25
	BArray< Cell_Type, Data_Type >, 45	counter_maxfuns, 26
	BArrayCell< Cell_Type, Data_Type >, 51	counter_neofun, 26
	BArrayDense < Cell_Type, Data_Type >, 64	counter_neofun_a2b, 26
	BArrayDenseCell< Cell_Type, Data_Type >, 71	counter_overall_changes, 26
	BArrayRow< Cell_Type, Data_Type >, 84	counter_overall_gains, 27
	BArrayVector < Cell_Type, Data_Type >, 89	counter_overall_gains_from_0, 27
oper	rator=	counter_overall_loss, 27
	BArray< Cell_Type, Data_Type >, 45	counter_pairwise_first_gain, 27
	BArrayCell< Cell_Type, Data_Type >, 51	counter_pairwise_neofun_singlefun, 28
	BArrayDense < Cell_Type, Data_Type >, 64	counter_pairwise_overall_change, 28
	BArrayDenseCell< Cell_Type, Data_Type >, 71	counter_pairwise_preserving, 28
	BArrayRow< Cell_Type, Data_Type >, 84	counter_preserve_pseudogene, 28
	BArrayVector< Cell_Type, Data_Type >, 90	counter_prop_genes_changing, 29
	Cell< Cell_Type >, 97, 98	counter_subfun, 29
	Counter< Array_Type, Data_Type >, 104	Phylo rules, 29
	Counters< Array_Type, Data_Type >, 108, 109	rule_dyn_limit_changes, 30
	Geese, 140, 141	phylo.hpp
	Model< Array_Type, Data_Counter_Type,	DEFAULT_DUPLICATION, 311
	Data Rule Type, Data Rule Dyn Type >,	DUPL_DUPL, 311
	158	DUPL_EITH, 311
	Rules< Array Type, Data Type >, 190	DUPL SPEC, 311
onoi		get_last_name, 316
opei	rator== PArray < Call Type Data Type > 45	IF MATCHES, 312
	BArray Cell Type, Data_Type >, 45	IF_NOTMATCHES, 312
	BArrayCell Cell_Type, Data_Type >, 51	IS_DUPLICATION, 312
	BArrayCell_const< Cell_Type, Data_Type >, 53	IS EITHER, 312
	BArrayDense < Cell_Type, Data_Type >, 64	IS_SPECIATION, 312
	BArrayDenseCell< Cell_Type, Data_Type >, 72	MAKE_DUPL_VARS, 313
	BArrayRow < Cell_Type, Data_Type >, 84	PHYLO_CHECK_MISSING, 313
	BArrayRow_const< Cell_Type, Data_Type >, 86	PHYLO_COUNTER_LAMBDA, 313
	BArrayVector< Cell_Type, Data_Type >, 90	PHYLO_RULE_DYN_LAMBDA, 313
	BArrayVector_const< Cell_Type, Data_Type >, 93	PhyloArray, 314
		i iiyioaiiay, o i -

PhyloCounter, 314	PhyloStatsCounter
PhyloCounters, 314	phylo.hpp, 315
PhyloModel, 314	PhyloSupport
PhyloPowerSet, 314	phylo.hpp, 316
PhyloRule, 315	POS
PhyloRuleData, 315	barraydense-meat-operators.hpp, 237
PhyloRuleDyn, 315	barraydense-meat.hpp, 242
PhyloRules, 315	barraydensecell-bones.hpp, 260
PhyloRulesDyn, 315	barraydensecell-meat.hpp, 262
PhyloStatsCounter, 315	barraydensecol-bones.hpp, 263
PhyloSupport, 316	barraydenserow-bones.hpp, 264
PHYLO_CHECK_MISSING	pos
	•
phylo COLNTED LAMBDA	PhyloRuleDynData, 176
PHYLO_COUNTER_LAMBDA	POS_N
phylo.hpp, 313	barraydense-meat-operators.hpp, 237
PHYLO_RULE_DYN_LAMBDA	barraydense-meat.hpp, 242
phylo.hpp, 313	barraydensecol-bones.hpp, 263
PhyloArray	barraydenserow-bones.hpp, 264
phylo.hpp, 314	PowerSet
PhyloCounter	PowerSet < Array_Type, Data_Rule_Type >, 179
phylo.hpp, 314	PowerSet < Array_Type, Data_Rule_Type >, 177
PhyloCounterData, 172	\sim PowerSet, 179
at, 173	add_rule, 179, 180
begin, 173	begin, 180
empty, 173	calc, 180
end, 174	coordinates_free, 182
get_counters, 174	coordinates_locked, 182
operator(), 174	data, 182
operator(), 174	EmptyArray, 182
·	
PhyloCounterData, 173	end, 180
push_back, 174	get_data, 180
reserve, 174	get_data_ptr, 181
shrink_to_fit, 175	init_support, 181
size, 175	M, 182
PhyloCounters	N, 183
phylo.hpp, 314	n_free, 183
PhyloModel	n_locked, 183
phylo.hpp, 314	operator[], 181
PhyloPowerSet	PowerSet, 179
phylo.hpp, 314	reset, 181
PhyloRule	rules, 183
phylo.hpp, 315	rules_deleted, 183
PhyloRuleData	size, 181
phylo.hpp, 315	predict
PhyloRuleDyn	Geese, 141
phylo.hpp, 315	predict_backend
PhyloRuleDynData, 175	Geese, 142
~PhyloRuleDynData, 176	
	predict_exhaust
counts, 176	Geese, 142
duplication, 176	predict_exhaust_backend
lb, 176	Geese, 142, 143
PhyloRuleDynData, 176	predict_sim
pos, 176	Geese, 143
ub, 177	print
PhyloRules	BArray< Cell_Type, Data_Type >, 45
phylo.hpp, 315	BArrayDense < Cell_Type, Data_Type >, 65
PhyloRulesDyn	Flock, 123
phylo.hpp, 315	FreqTable< T >, 127
	•

Geese, 143	counters-meat.hpp, 289
Model < Array_Type, Data_Counter_Type,	statscounter-meat.hpp, 343
Data_Rule_Type, Data_Rule_Dyn_Type >,	support-meat.hpp, 354
158	rhs
Support < Array_Type, Data_Counter_Type,	barray-meat-operators.hpp, 213
Data_Rule_Type, Data_Rule_Dyn_Type >, 200	rm_cell BArray< Cell_Type, Data_Type >, 46
print_observed_counts	BArrayDense< Cell_Type, Data_Type >, 40
Geese, 144	barraydense-meat.hpp, 253
print_stats	support-meat.hpp, 349
Model< Array_Type, Data_Counter_Type,	ROW
Data_Rule_Type, Data_Rule_Dyn_Type >,	barray-meat-operators.hpp, 211
158	barray-meat.hpp, 217, 225, 226
printf_barry	barraydense-meat-operators.hpp, 237
barry-configuration.hpp, 271	barraydense-meat.hpp, 242
probability	row
Node, 170	BArray< Cell_Type, Data_Type >, 46
Progress, 184	BArrayDense< Cell_Type, Data_Type >, 65, 66
\sim Progress, 184	row0
end, 185	barray-meat.hpp, 231
next, 185	Row_type
Progress, 184	typedefs.hpp, 358
progress.hpp	rowsum
BARRY_PROGRESS_BAR_WIDTH, 334	BArrayDense < Cell_Type, Data_Type >, 66
pset_loc	Rule
Geese, 146	Rule < Array_Type, Data_Type >, 186
push_back PhyloCounterData 174	Rule< Array_Type, Data_Type >, 185 ~Rule, 186
PhyloCounterData, 174	D, 187
README.md, 360	operator(), 187
reduced_sequence	Rule, 186
Geese, 147	rule_dyn_limit_changes
rengine	Phylo rules, 30
Flock, 124	rule_fun_default
report	rules-bones.hpp, 335
barray-meat.hpp, 230	Rule_fun_type
barraydense-meat.hpp, 258	typedefs.hpp, 359
reserve	RULE_FUNCTION
BArray < Cell_Type, Data_Type >, 46	barry.hpp, 276
BArrayDense < Cell_Type, Data_Type >, 65	defm-bones.hpp, 322
FreqTable < T >, 127	geese-bones.hpp, 326
PhyloCounterData, 174	RULE_LAMBDA
reset PowerCet < Array Time Deta Dula Time > 101	barry.hpp, 276
PowerSet < Array_Type, Data_Rule_Type >, 181	Rules
reset_array StatsCounter< Array_Type, Data_Type >, 193	Rules < Array_Type, Data_Type >, 188
Support< Array_Type, Data_Type >, 100	rules
Data_Rule_Type, Data_Rule_Dyn_Type >,	PowerSet < Array_Type, Data_Rule_Type >, 183
200, 201	support-meat.hpp, 355
resize	Rules < Array_Type, Data_Type >, 187
BArray< Cell_Type, Data_Type >, 46	~Rules, 188
barray-meat.hpp, 225	add_rule, 189 get_seq, 189
BArrayDense< Cell_Type, Data_Type >, 65	operator(), 190
barraydense-meat.hpp, 252	operator=, 190
Entries < Cell_Type >, 117	Rules, 188
statscounter-meat.hpp, 340	size, 190
return	rules-bones.hpp
barray-meat.hpp, 225, 230	rule_fun_default, 335
barraydense-meat.hpp, 258	,

rules_	Model < Array_Type, Data_Counter_Type,
support-meat.hpp, 355	Data_Rule_Type, Data_Rule_Dyn_Type >,
rules_deleted	160
PowerSet< Array_Type, Data_Rule_Type >, 183	shrink_to_fit
rules_dyn	PhyloCounterData, 175
support-meat.hpp, 355	simulate
rules_zerodiag	Geese, 144
DEFMArray counters, 21	size
network.hpp, 308	BArrayDenseCol< Cell_Type, Data_Type >, 74
networkinpp, ooo	BArrayDenseCol_const< Cell_Type, Data_Type >,
sample	77
Model< Array_Type, Data_Counter_Type,	
Data_Rule_Type, Data_Rule_Dyn_Type >,	BArrayDenseRow< Cell_Type, Data_Type >, 79
158, 159	BArrayDenseRow_const< Cell_Type, Data_Type
	>, 81
search	BArrayVector< Cell_Type, Data_Type >, 90
barray-meat.hpp, 231	BArrayVector_const< Cell_Type, Data_Type >, 94
sequence	Counters< Array_Type, Data_Type >, 109
Geese, 147	FreqTable $<$ T $>$, 127
set_counters	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 >,	161
159	PhyloCounterData, 175
StatsCounter< Array_Type, Data_Type >, 194	PowerSet < Array_Type, Data_Rule_Type >, 181
Support< Array_Type, Data_Counter_Type,	Rules < Array_Type, Data_Type >, 190
Data_Rule_Type, Data_Rule_Dyn_Type >,	StatsCounter< Array_Type, Data_Type >, 194
201	size_unique
set_data	Model< Array_Type, Data_Counter_Type,
BArray< Cell_Type, Data_Type >, 46	Data_Rule_Type, Data_Rule_Dyn_Type >,
BArrayDense < Cell_Type, Data_Type >, 66	161
set_keygen	source
Model< Array_Type, Data_Counter_Type,	barray-meat.hpp, 231
Data_Rule_Type, Data_Rule_Dyn_Type >,	barraydense-meat.hpp, 258
159	Entries < Cell_Type >, 117
set_rengine	states
Model< Array_Type, Data_Counter_Type,	
Data_Rule_Type, Data_Rule_Dyn_Type >,	NodeData, 172
159	Statistical Models, 11
set rules	stats_bank
Model< Array_Type, Data_Counter_Type,	support-meat.hpp, 355
Data_Rule_Type, Data_Rule_Dyn_Type >,	StatsCounter
159	StatsCounter< Array_Type, Data_Type >, 191,
	192
Support< Array_Type, Data_Counter_Type,	StatsCounter< Array_Type, Data_Type >, 191
Data_Rule_Type, Data_Rule_Dyn_Type >,	\sim StatsCounter, 192
201	add_counter, 192
set_rules_dyn	count_all, 192
Model< Array_Type, Data_Counter_Type,	count_current, 193
Data_Rule_Type, Data_Rule_Dyn_Type >,	count_init, 193
160	get_counters, 193
Support< Array_Type, Data_Counter_Type,	get_descriptions, 193
Data_Rule_Type, Data_Rule_Dyn_Type >,	get_names, 193
201	reset_array, 193
set_seed	set_counters, 194
Flock, 123	size, 194
Geese, 144	StatsCounter, 191, 192
Model < Array_Type, Data_Counter_Type,	statscounter-meat.hpp
Data_Rule_Type, Data_Rule_Dyn_Type >,	clear, 340
160	counter, 342
set_transform_model	counter_deleted, 342
	Countei_ucicteu, 342

counters, 342	print, 200
counters_, 342	reset_array, 200, 201
current_stats, 342	set_counters, 201
EmptyArray, 343	set_rules, 201
f_, 343	set_rules_dyn, 201
for, 340	Support, 196, 197
j, 343	support-meat.hpp
resize, 340	array_bank, 352
return, 343	BARRY_SUPPORT_MEAT_HPP, 347
STATSCOUNTER_TEMPLATE, 339–341	calc_backend_dense, 347
STATSCOUNTER_TEMPLATE_ARGS, 339	calc_backend_sparse, 348
STATSCOUNTER_TYPE, 339 STATSCOUNTER_TEMPLATE	change_stats_different, 352
	coord_i, 353
statscounter-meat.hpp, 339–341 STATSCOUNTER_TEMPLATE_ARGS	coord_j, 353 counters, 353
statscounter-meat.hpp, 339	counters_, 353
STATSCOUNTER_TYPE	delete_counters, 353
statscounter-meat.hpp, 339	delete_counters, 353
store_psets	delete_rules_dyn, 354
Model< Array_Type, Data_Counter_Type,	else, 354
Data_Rule_Type, Data_Rule_Dyn_Type >,	f_, 354
161	for, 348
subtree_prob	hashes, 354
Node, 170	if, 348
Support	insert_cell, 348, 349
Support< Array_Type, Data_Counter_Type,	return, 354
Data_Rule_Type, Data_Rule_Dyn_Type >,	rm_cell, 349
196, 197	rules, 355
Support < Array_Type, Data_Counter_Type, Data_Rule_1	
Data_Rule_Dyn_Type >, 194	rules_dyn, 355
\sim Support, 197	stats_bank, 355
add_counter, 197	SUPPORT_TEMPLATE, 347, 349-352
add_rule, 197, 198	SUPPORT_TEMPLATE_ARGS, 347
add_rule_dyn, 198	SUPPORT_TYPE, 347
calc, 198	tmp_chng, 355
change_stats, 201	support_size
coordiantes_n_free, 202	Flock, 123
coordiantes_n_locked, 202	Geese, 145
coordinates_free, 202	Model < Array_Type, Data_Counter_Type,
coordinates_locked, 202	Data_Rule_Type, Data_Rule_Dyn_Type >,
current_stats, 202	161
delete_counters, 203	SUPPORT_TEMPLATE
delete_rules, 203	support-meat.hpp, 347, 349-352
delete_rules_dyn, 203	SUPPORT_TEMPLATE_ARGS
eval_rules_dyn, 199	support-meat.hpp, 347
get_counters, 199	SUPPORT_TYPE
get_counts, 199	support-meat.hpp, 347
get_current_stats, 199	swap_cells
get_data, 199	BArray< Cell_Type, Data_Type >, 47
get_rules, 200	BArrayDense< Cell_Type, Data_Type >, 66
get_rules_dyn, 200	swap_cols
hashes, 203	BArray Cell_Type, Data_Type >, 47
hashes_initialized, 203	BArrayDense < Cell_Type, Data_Type >, 67
init_support, 200	swap_rows PArroy < Coll Type Date Type > 47
M, 204	BArray Cell_Type, Data_Type >, 47
max_num_elements, 204	BArrayDense< Cell_Type, Data_Type >, 67
N, 204 n_counters, 204	target
ii_counters, 204	barray-meat.hpp, 231
	1.1.1 -

barraydense-meat.hpp, 259	value
Entries < Cell_Type >, 117	barray-meat.hpp, 231
this	barraydense-meat.hpp, 259
barray-meat-operators.hpp, 213	Cell< Cell_Type >, 98
counters-meat.hpp, 289	vec_diff
tmp_chng	defm-bones.hpp, 322
support-meat.hpp, 355	geese-bones.hpp, 326
toggle_cell	vec_equal
BArray< Cell_Type, Data_Type >, 47	typedefs.hpp, 359
BArrayDense < Cell_Type, Data_Type >, 67	vec_equal_approx
toggle_lock	typedefs.hpp, 359
BArray< Cell_Type, Data_Type >, 47	vec_inner_prod
BArrayDense< Cell_Type, Data_Type >, 67	typedefs.hpp, 360
transform_model	vecHasher< T >, 205
Model< Array_Type, Data_Counter_Type,	operator(), 205
Data_Rule_Type, Data_Rule_Dyn_Type >,	vector_caster
161	geese-bones.hpp, 326
transpose	vertex_attr
BArray< Cell_Type, Data_Type >, 48	NetworkData, 165
BArrayDense < Cell_Type, Data_Type >, 40	visited
TWO	BArray< Cell_Type, Data_Type >, 49
CHECK, 33	BArrayDense< Cell_Type, Data_Type >, 49
EXISTS, 34	
	Cell< Cell_Type >, 98
typedefs.hpp	Node, 170
Col_type, 357	vprintf
Counter_fun_type, 358	barraydense-meat.hpp, 253
Counts_type, 358	ZERO_CELL
MapVec_type, 358	barraydense-meat.hpp, 243
Row_type, 358	barraydensecol-bones.hpp, 263
Rule_fun_type, 359	barraydenserow-bones.hpp, 264
uint, 359	
vec_equal, 359	zero_col
vec_equal_approx, 359	BArray Cell_Type, Data_Type >, 48
vec_inner_prod, 360	BArrayDense < Cell_Type, Data_Type >, 68
la	Zero_row
Ub	BArray Cell_Type, Data_Type >, 48
PhyloRuleDynData, 177	BArrayDense< Cell_Type, Data_Type >, 68
uint	
typedefs.hpp, 359 UKNOWN	
EXISTS, 34	
update_annotations	
Geese, 145	
update_normalizing_constant	
model-meat.hpp, 321	
V	
barray-meat.hpp, 231	
barraydense-meat.hpp, 259	
• • • • • • • • • • • • • • • • • • • •	
va_end	
barraydense-meat.hpp, 253	
va_start	
barraydense-meat.hpp, 253	
Entries < Cell_Type >, 117	
val0	
barraydense-meat.hpp, 259	
val1	
barraydense-meat.hpp, 259	
parrayuerise-ineal.hpp, 200	