barry: Your go-to motif accountant 0.0-1

Generated by Doxygen 1.9.1

1 Main Page	1
2 Module Index	3
2.1 Modules	3
3 Class Index	5
3.1 Class List	5
4 File Index	7
4.1 File List	7
5 Module Documentation	9
5.1 Counting	9
•	9
·	9
5.2.1 Detailed Description	0
5.3 Network counters	0
5.3.1 Detailed Description	1
5.3.2 Function Documentation	1
5.3.2.1 counter_absdiff()	1
5.3.2.2 counter_css_partially_false_recip()	1
5.3.2.3 counter_ctriads()	1
5.3.2.4 counter_degree()	2
5.3.2.5 counter_density()	2
5.3.2.6 counter_diff()	2
5.3.2.7 counter_edges()	2
5.3.2.8 counter_idegree()	3
5.3.2.9 counter_idegree15()	3
5.3.2.10 counter_isolates()	3
5.3.2.11 counter_istar2()	3
5.3.2.12 counter_mutual()	3
5.3.2.13 counter_nodecov()	4
5.3.2.14 counter_nodeicov()	4
5.3.2.15 counter_nodematch()	4
5.3.2.16 counter_nodeocov()	4
5.3.2.17 counter_odegree()	4
5.3.2.18 counter_odegree15()	5
5.3.2.19 counter_ostar2()	5
5.3.2.20 counter_ttriads()	5
5.3.2.21 NETWORK_COUNTER()	5
5.4 Phylo counters	5
5.4.1 Detailed Description	6
5.4.2 Function Documentation	6
5.4.2.1 counter_co_opt()	7

	5.4.2.2 counter_cogain()	17
	5.4.2.3 counter_gains()	17
	5.4.2.4 counter_gains_k_offspring()	18
	5.4.2.5 counter_genes_changing()	18
	5.4.2.6 counter_longest()	18
	5.4.2.7 counter_loss()	18
	5.4.2.8 counter_maxfuns()	19
	5.4.2.9 counter_neofun()	19
	5.4.2.10 counter_neofun_a2b()	19
	5.4.2.11 counter_overall_changes()	19
	5.4.2.12 counter_overall_gains()	20
	5.4.2.13 counter_overall_loss()	20
	5.4.2.14 counter_prop_genes_changing()	20
	5.4.2.15 counter_subfun()	20
	5.5 Phylo rules	21
	5.5.1 Detailed Description	21
	5.5.2 Function Documentation	21
	5.5.2.1 rule_dyn_limit_changes()	21
6	Namespace Documentation	23
0	6.1 barry Namespace Reference	23 23
	6.1.1 Detailed Description	
	6.2 barry::counters Namespace Reference	23
	6.2.1 Detailed Description	23
	6.3 barry::counters::network Namespace Reference	23
	6.4 barry::counters::phylo Namespace Reference	24
	6.5 CHECK Namespace Reference	24
	6.5.1 Detailed Description	
	6.5.2 Variable Documentation	24 24
	6.5.2.1 BOTH	24
	6.5.2.2 NONE	24 24
	6.5.2.3 ONE	24
	6.5.2.4 TWO	25
	6.6 EXISTS Namespace Reference	25
	6.6.1 Detailed Description	25
	6.6.2 Variable Documentation	25
	6.6.2.1 AS_ONE	25
	6.6.2.2 AS_ZERO	25
	6.6.2.3 BOTH	25 26
	6.6.2.4 NONE	26 26
	6.6.2.5 ONE	26 26
	6.6.2.6 TWO	
	U.U.C.U 1 ** U	20

6.6.2.7 UKNOWN	. 26
7 Class Documentation	27
7.1 BArray< Cell_Type, Data_Type > Class Template Reference	. 27
7.1.1 Detailed Description	. 29
7.1.2 Constructor & Destructor Documentation	. 30
7.1.2.1 BArray() [1/6]	. 30
7.1.2.2 BArray() [2/6]	. 30
7.1.2.3 BArray() [3/6]	. 30
7.1.2.4 BArray() [4/6]	. 31
7.1.2.5 BArray() [5/6]	. 31
7.1.2.6 BArray() [6/6]	. 31
7.1.2.7 ~BArray()	. 31
7.1.3 Member Function Documentation	. 31
7.1.3.1 clear()	. 31
7.1.3.2 col()	. 32
7.1.3.3 D() [1/2]	. 32
7.1.3.4 D() [2/2]	. 32
7.1.3.5 default_val()	. 32
7.1.3.6 flush_data()	. 32
7.1.3.7 get_cell()	. 32
7.1.3.8 get_col_vec() [1/2]	. 33
7.1.3.9 get_col_vec() [2/2]	. 33
7.1.3.10 get_entries()	. 33
7.1.3.11 get_row_vec() [1/2]	. 33
7.1.3.12 get_row_vec() [2/2]	. 33
7.1.3.13 insert_cell() [1/3]	. 34
7.1.3.14 insert_cell() [2/3]	. 34
7.1.3.15 insert_cell() [3/3]	. 34
7.1.3.16 is_empty()	. 34
7.1.3.17 ncol()	. 34
7.1.3.18 nnozero()	. 35
7.1.3.19 nrow()	. 35
7.1.3.20 operator()() [1/2]	. 35
7.1.3.21 operator()() [2/2]	. 35
7.1.3.22 operator*=()	. 35
7.1.3.23 operator+=() [1/3]	. 35
7.1.3.24 operator+=() [2/3]	. 36
7.1.3.25 operator+=() [3/3]	. 36
7.1.3.26 operator-=() [1/3]	. 36
7.1.3.27 operator-=() [2/3]	. 36
7.1.3.28 operator-=() [3/3]	. 36

7.1.3.29 operator/=()	36
7.1.3.30 operator=() [1/2]	37
7.1.3.31 operator=() [2/2]	37
7.1.3.32 operator==()	37
7.1.3.33 out_of_range()	37
7.1.3.34 print()	37
7.1.3.35 reserve()	37
7.1.3.36 resize()	38
7.1.3.37 rm_cell()	38
7.1.3.38 row()	38
7.1.3.39 set_data()	38
7.1.3.40 swap_cells()	38
7.1.3.41 swap_cols()	39
7.1.3.42 swap_rows()	39
7.1.3.43 toggle_cell()	39
7.1.3.44 toggle_lock()	39
7.1.3.45 transpose()	40
7.1.3.46 zero_col()	40
7.1.3.47 zero_row()	40
7.1.4 Friends And Related Function Documentation	40
7.1.4.1 BArrayCell< Cell_Type, Data_Type >	40
7.1.4.2 BArrayCell_const< Cell_Type, Data_Type >	40
7.1.5 Member Data Documentation	40
7.1.5.1 visited	41
7.2 BArrayCell< Cell_Type, Data_Type > Class Template Reference	41
7.2.1 Detailed Description	41
7.2.2 Constructor & Destructor Documentation	41
7.2.2.1 BArrayCell()	42
7.2.2.2 ~BArrayCell()	42
7.2.3 Member Function Documentation	42
7.2.3.1 operator Cell_Type()	42
7.2.3.2 operator*=()	42
7.2.3.3 operator+=()	42
7.2.3.4 operator-=()	43
7.2.3.5 operator/=()	43
7.2.3.6 operator=()	43
7.2.3.7 operator==()	43
7.3 BArrayCell_const< Cell_Type, Data_Type > Class Template Reference	43
7.3.1 Detailed Description	44
7.3.2 Constructor & Destructor Documentation	44
7.3.2.1 BArrayCell_const()	44
7.3.2.2 ~BArrayCell_const()	44

7.3.3 Member Function Documentation	44
7.3.3.1 operator Cell_Type()	45
7.3.3.2 operator"!=()	45
7.3.3.3 operator<()	45
7.3.3.4 operator<=()	45
7.3.3.5 operator==()	45
7.3.3.6 operator>()	46
7.3.3.7 operator>=()	46
7.4 BArrayDense< Cell_Type, Data_Type > Class Template Reference	46
7.4.1 Detailed Description	48
7.4.2 Constructor & Destructor Documentation	49
7.4.2.1 BArrayDense() [1/6]	49
7.4.2.2 BArrayDense() [2/6]	49
7.4.2.3 BArrayDense() [3/6]	49
7.4.2.4 BArrayDense() [4/6]	50
7.4.2.5 BArrayDense() [5/6]	50
7.4.2.6 BArrayDense() [6/6]	50
7.4.2.7 ~BArrayDense()	50
7.4.3 Member Function Documentation	50
7.4.3.1 clear()	51
7.4.3.2 col()	51
7.4.3.3 D() [1/2]	51
7.4.3.4 D() [2/2]	51
7.4.3.5 default_val()	51
7.4.3.6 get_cell()	52
7.4.3.7 get_col_vec() [1/2]	52
7.4.3.8 get_col_vec() [2/2]	52
7.4.3.9 get_entries()	52
7.4.3.10 get_row_vec() [1/2]	53
7.4.3.11 get_row_vec() [2/2]	53
7.4.3.12 insert_cell() [1/3]	53
7.4.3.13 insert_cell() [2/3]	53
7.4.3.14 insert_cell() [3/3]	54
7.4.3.15 is_empty()	54
7.4.3.16 ncol()	54
7.4.3.17 nnozero()	54
7.4.3.18 nrow()	54
7.4.3.19 operator()() [1/2]	55
7.4.3.20 operator()() [2/2]	55
7.4.3.21 operator*=()	55
7.4.3.22 operator+=() [1/3]	55
7.4.3.23 operator+=() [2/3]	55

7.4.3.24 operator+=() [3/3]	20
7.4.3.25 operator-=() [1/3]	56
7.4.3.26 operator-=() [2/3]	56
7.4.3.27 operator-=() [3/3]	56
7.4.3.28 operator/=()	56
7.4.3.29 operator=() [1/2]	57
7.4.3.30 operator=() [2/2]	57
7.4.3.31 operator==()	57
7.4.3.32 out_of_range()	57
7.4.3.33 print()	57
7.4.3.34 reserve()	58
7.4.3.35 resize()	58
7.4.3.36 rm_cell()	58
7.4.3.37 row()	58
7.4.3.38 set_data()	58
7.4.3.39 swap_cells()	59
7.4.3.40 swap_cols()	59
7.4.3.41 swap_rows()	59
7.4.3.42 toggle_cell()	60
7.4.3.43 toggle_lock()	60
7.4.3.44 transpose()	60
7.4.3.45 zero_col()	60
7.4.3.46 zero_row()	60
7.4.4 Friends And Related Function Documentation	61
7.4.4.1 BArrayCell< Cell_Type, Data_Type >	61
7.4.4.2 BArrayCell_const< Cell_Type, Data_Type >	61
7.4.5 Member Data Documentation	61
7.4.5.1 visited	61
7.5 BArrayDenseCell < Cell_Type, Data_Type > Class Template Reference	61
7.5.1 Detailed Description	62
7.5.2 Constructor & Destructor Documentation	62
7.5.2.1 BArrayDenseCell()	62
7.5.2.2 ~BArrayDenseCell()	62
7.5.3 Member Function Documentation	62
7.5.3.1 operator Cell_Type()	63
7.5.3.2 operator*=()	63
7.5.3.3 operator+=()	63
7.5.3.4 operator-=()	63
7.5.3.5 operator/=()	63
7.5.3.6 operator=()	64
7.5.3.7 operator==()	64
7.6 BArrayDenseCell_const< Cell_Type, Data_Type > Class Template Reference	64

7.6.1 Detailed Description	64
7.6.2 Constructor & Destructor Documentation	65
7.6.2.1 BArrayDenseCell_const()	65
$7.6.2.2 \sim BArrayDenseCell_const() \ . \ . \ . \ . \ . \ . \ . \ . \ . \ $	65
7.6.3 Member Function Documentation	65
7.6.3.1 operator Cell_Type()	65
7.6.3.2 operator"!=()	65
7.6.3.3 operator<()	66
7.6.3.4 operator<=()	66
7.6.3.5 operator==()	66
7.6.3.6 operator>()	66
7.6.3.7 operator>=()	66
7.7 BArrayVector< Cell_Type, Data_Type > Class Template Reference	67
7.7.1 Detailed Description	67
7.7.2 Constructor & Destructor Documentation	67
7.7.2.1 BArrayVector()	67
7.7.2.2 ~BArrayVector()	68
7.7.3 Member Function Documentation	68
7.7.3.1 begin()	68
7.7.3.2 end()	68
7.7.3.3 is_col()	69
7.7.3.4 is_row()	69
7.7.3.5 operator std::vector< Cell_Type >()	69
7.7.3.6 operator*=()	69
7.7.3.7 operator+=()	69
7.7.3.8 operator-=()	70
7.7.3.9 operator/=()	70
7.7.3.10 operator=()	70
7.7.3.11 operator==()	70
7.7.3.12 size()	70
$7.8\; BArray Vector_const < Cell_Type, \; Data_Type > Class\; Template\; Reference \\ \; \ldots \\ \;$	71
7.8.1 Detailed Description	71
7.8.2 Constructor & Destructor Documentation	71
7.8.2.1 BArrayVector_const()	71
$7.8.2.2 \sim BArrayVector_const() \ . \ . \ . \ . \ . \ . \ . \ . \ . \ $	72
7.8.3 Member Function Documentation	72
7.8.3.1 begin()	72
7.8.3.2 end()	72
7.8.3.3 is_col()	72
7.8.3.4 is_row()	72
7.8.3.5 operator std::vector $<$ Cell_Type $>$ ()	72
7.8.3.6 operator"!=()	73

7.8.3.7 operator<()	. 73
7.8.3.8 operator<=()	. 73
7.8.3.9 operator==()	. 73
7.8.3.10 operator>()	. 73
7.8.3.11 operator>=()	. 74
7.8.3.12 size()	. 74
7.9 Cell< Cell_Type > Class Template Reference	. 74
7.9.1 Detailed Description	. 75
7.9.2 Constructor & Destructor Documentation	. 75
7.9.2.1 Cell() [1/7]	. 75
7.9.2.2 Cell() [2/7]	. 75
7.9.2.3 ~Cell()	. 75
7.9.2.4 Cell() [3/7]	. 76
7.9.2.5 Cell() [4/7]	. 76
7.9.2.6 Cell() [5/7]	. 76
7.9.2.7 Cell() [6/7]	. 76
7.9.2.8 Cell() [7/7]	. 76
7.9.3 Member Function Documentation	. 76
7.9.3.1 add() [1/4]	. 77
7.9.3.2 add() [2/4]	. 77
7.9.3.3 add() [3/4]	. 77
7.9.3.4 add() [4/4]	. 77
7.9.3.5 operator Cell_Type()	. 77
7.9.3.6 operator"!=()	. 77
7.9.3.7 operator=() [1/2]	. 78
7.9.3.8 operator=() [2/2]	. 78
7.9.3.9 operator==()	. 78
7.9.4 Member Data Documentation	. 78
7.9.4.1 value	. 78
7.9.4.2 visited	. 78
7.10 ConstBArrayRowlter< Cell_Type, Data_Type > Class Template Reference	. 79
7.10.1 Detailed Description	. 79
7.10.2 Constructor & Destructor Documentation	. 79
7.10.2.1 ConstBArrayRowlter()	. 80
7.10.2.2 ~ConstBArrayRowlter()	. 80
7.10.3 Member Data Documentation	. 80
7.10.3.1 Array	. 80
7.10.3.2 current_col	. 80
7.10.3.3 current_row	. 80
7.10.3.4 iter	. 81
7.11 Counter< Array_Type, Data_Type > Class Template Reference	. 81
7.11.1 Detailed Description	82

7.11.2 Constructor & Destructor Documentation	82
7.11.2.1 Counter() [1/4]	82
7.11.2.2 Counter() [2/4]	82
7.11.2.3 Counter() [3/4]	83
7.11.2.4 Counter() [4/4]	83
7.11.2.5 ~Counter()	83
7.11.3 Member Function Documentation	83
7.11.3.1 count()	83
7.11.3.2 init()	84
7.11.3.3 operator=() [1/2]	84
7.11.3.4 operator=() [2/2]	84
7.11.4 Member Data Documentation	84
7.11.4.1 count_fun	84
7.11.4.2 data	85
7.11.4.3 delete_data	85
7.11.4.4 desc	85
7.11.4.5 init_fun	85
7.11.4.6 name	85
7.12 Counters< Array_Type, Data_Type > Class Template Reference	86
7.12.1 Detailed Description	86
7.12.2 Constructor & Destructor Documentation	86
7.12.2.1 Counters() [1/3]	87
7.12.2.2 ~Counters()	87
7.12.2.3 Counters() [2/3]	87
7.12.2.4 Counters() [3/3]	87
7.12.3 Member Function Documentation	88
7.12.3.1 add_counter() [1/3]	88
7.12.3.2 add_counter() [2/3]	88
7.12.3.3 add_counter() [3/3]	88
7.12.3.4 clear()	88
7.12.3.5 operator=() [1/2]	88
7.12.3.6 operator=() [2/2]	89
7.12.3.7 operator[]()	89
7.12.3.8 size()	90
7.13 Entries $<$ Cell_Type $>$ Class Template Reference	90
7.13.1 Detailed Description	90
7.13.2 Constructor & Destructor Documentation	91
7.13.2.1 Entries() [1/2]	91
7.13.2.2 Entries() [2/2]	91
7.13.2.3 ~Entries()	91
7.13.3 Member Function Documentation	91
7.13.3.1 resize()	91

7.13.4 Member Data Documentation	92
7.13.4.1 source	92
7.13.4.2 target	92
7.13.4.3 val	92
7.14 Flock Class Reference	92
7.14.1 Detailed Description	93
7.14.2 Constructor & Destructor Documentation	93
7.14.2.1 Flock()	94
7.14.2.2 ~Flock()	94
7.14.3 Member Function Documentation	94
7.14.3.1 add_data()	94
7.14.3.2 colnames()	94
7.14.3.3 get_counters()	95
7.14.3.4 get_model()	95
7.14.3.5 get_support()	95
7.14.3.6 init()	95
7.14.3.7 likelihood_joint()	95
7.14.3.8 nfuns()	96
7.14.3.9 nleafs()	96
7.14.3.10 nnodes()	96
7.14.3.11 nterms()	96
7.14.3.12 ntrees()	96
7.14.3.13 operator()()	96
7.14.3.14 parse_polytomies()	97
7.14.3.15 print()	97
7.14.3.16 set_seed()	97
7.14.3.17 support_size()	97
7.14.4 Member Data Documentation	98
7.14.4.1 dat	98
7.14.4.2 initialized	98
7.14.4.3 model	98
7.14.4.4 nfunctions	98
7.14.4.5 rengine	98
7.15 FreqTable < T > Class Template Reference	99
7.15.1 Detailed Description	99
7.15.2 Constructor & Destructor Documentation	99
7.15.2.1 FreqTable()	99
7.15.2.2 ~FreqTable()	99
7.15.3 Member Function Documentation	100
7.15.3.1 add()	100
7.15.3.2 as_vector()	100
7.15.3.3 clear()	100

7.15.3.4 get_data()	100
7.15.3.5 get_data_ptr()	100
7.15.3.6 print()	101
7.15.3.7 reserve()	101
7.15.3.8 size()	101
7.16 Geese Class Reference	101
7.16.1 Detailed Description	104
7.16.2 Constructor & Destructor Documentation	104
7.16.2.1 Geese() [1/4]	104
7.16.2.2 Geese() [2/4]	104
7.16.2.3 Geese() [3/4]	104
7.16.2.4 Geese() [4/4]	105
7.16.2.5 ~Geese()	105
7.16.3 Member Function Documentation	105
7.16.3.1 calc_reduced_sequence()	105
7.16.3.2 calc_sequence()	105
7.16.3.3 colnames()	105
7.16.3.4 get_annotated_nodes()	106
7.16.3.5 get_counters()	106
7.16.3.6 get_model()	106
7.16.3.7 get_probabilities()	106
7.16.3.8 get_rengine()	106
7.16.3.9 get_states()	107
7.16.3.10 get_support()	107
7.16.3.11 inherit_support()	107
7.16.3.12 init()	107
7.16.3.13 init_node()	107
7.16.3.14 likelihood()	108
7.16.3.15 likelihood_exhaust()	108
7.16.3.16 nannotations()	108
7.16.3.17 nfuns()	108
7.16.3.18 nleafs()	108
7.16.3.19 nnodes()	109
7.16.3.20 nterms()	109
7.16.3.21 observed_counts()	109
7.16.3.22 operator=() [1/2]	109
7.16.3.23 operator=() [2/2]	109
7.16.3.24 parse_polytomies()	109
7.16.3.25 predict()	110
7.16.3.26 predict_backend()	110
7.16.3.27 predict_exhaust()	110
7.16.3.28 predict_exhaust_backend()	110

7.16.3.29 predict_sim()	
7.16.3.30 print()	111
7.16.3.31 print_observed_counts()	111
7.16.3.32 set_seed()	111
7.16.3.33 simulate()	111
7.16.3.34 support_size()	111
7.16.3.35 update_annotations()	
7.16.4 Member Data Documentation	
7.16.4.1 delete_rengine	
7.16.4.2 delete_support	
7.16.4.3 initialized	
7.16.4.4 map_to_nodes	
7.16.4.5 nfunctions	
7.16.4.6 nodes	
7.16.4.7 reduced_sequence	
7.16.4.8 sequence	
7.17 Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule	
plate Reference	
7.17.1 Detailed Description	
7.17.2 Constructor & Destructor Documentation	
7.17.2.1 Model() [1/3]	
7.17.2.2 Model() [2/3]	
7.17.2.3 Model() [3/3]	
7.17.2.4 ~Model()	
7.17.3 Member Function Documentation	
7.17.3.1 add_array()	
7.17.3.2 add_counter() [1/3]	
7.17.3.3 add_counter() [2/3]	
7.17.3.4 add_counter() [3/3]	
7.17.3.5 add_rule() [1/3]	
7.17.3.6 add_rule() [2/3]	
7.17.3.7 add_rule() [3/3]	
7.17.3.8 add_rule_dyn() [1/3]	
7.17.3.9 add_rule_dyn() [2/3]	
7.17.3.10 add_rule_dyn() [3/3]	
7.17.3.11 colnames()	
7.17.3.12 conditional_prob()	
7.17.3.13 gen_key()	
7.17.3.14 get_counters()	121
7.17.3.15 get_norm_const()	121
7.17.3.16 get_pset()	121
7.17.3.17 get_pset_stats()	

7.17.3.18 get_rengine()	. 122
7.17.3.19 get_rules()	. 122
7.17.3.20 get_rules_dyn()	. 122
7.17.3.21 get_support()	. 122
7.17.3.22 likelihood() [1/3]	. 123
7.17.3.23 likelihood() [2/3]	. 123
7.17.3.24 likelihood() [3/3]	. 123
7.17.3.25 likelihood_total()	. 123
7.17.3.26 nterms()	. 124
7.17.3.27 operator=()	. 124
7.17.3.28 print()	. 124
7.17.3.29 print_stats()	. 124
7.17.3.30 sample() [1/2]	. 124
7.17.3.31 sample() [2/2]	. 125
7.17.3.32 set_counters()	. 125
7.17.3.33 set_keygen()	. 125
7.17.3.34 set_rengine()	. 125
7.17.3.35 set_rules()	. 125
7.17.3.36 set_rules_dyn()	. 126
7.17.3.37 set_seed()	. 126
7.17.3.38 size()	. 126
7.17.3.39 size_unique()	. 126
7.17.3.40 store_psets()	. 126
7.17.3.41 support_size()	. 127
7.18 NetCounterData Class Reference	. 127
7.18.1 Detailed Description	. 127
7.18.2 Constructor & Destructor Documentation	. 127
7.18.2.1 NetCounterData() [1/2]	. 127
7.18.2.2 NetCounterData() [2/2]	. 128
7.18.2.3 ∼NetCounterData()	. 128
7.18.3 Member Data Documentation	. 128
7.18.3.1 indices	. 128
7.18.3.2 numbers	. 128
7.19 NetworkData Class Reference	. 128
7.19.1 Detailed Description	. 129
7.19.2 Constructor & Destructor Documentation	. 129
7.19.2.1 NetworkData() [1/3]	. 129
7.19.2.2 NetworkData() [2/3]	. 129
7.19.2.3 NetworkData() [3/3]	. 130
7.19.2.4 ~NetworkData()	. 130
7.19.3 Member Data Documentation	. 130
7.19.3.1 directed	. 130

7.19.3.2 vertex_attr	131
7.20 Node Class Reference	131
7.20.1 Detailed Description	132
7.20.2 Constructor & Destructor Documentation	132
7.20.2.1 Node() [1/5]	132
7.20.2.2 Node() [2/5]	133
7.20.2.3 Node() [3/5]	133
7.20.2.4 Node() [4/5]	133
7.20.2.5 Node() [5/5]	133
7.20.2.6 ~Node()	133
7.20.3 Member Function Documentation	133
7.20.3.1 get_parent()	134
7.20.3.2 is_leaf()	134
7.20.3.3 noffspring()	134
7.20.4 Member Data Documentation	134
7.20.4.1 annotations	134
7.20.4.2 array	134
7.20.4.3 arrays	135
7.20.4.4 duplication	135
7.20.4.5 id	135
7.20.4.6 narray	135
7.20.4.7 offspring	135
7.20.4.8 ord	136
7.20.4.9 parent	136
7.20.4.10 probability	136
7.20.4.11 subtree_prob	136
7.20.4.12 visited	136
7.21 NodeData Class Reference	137
7.21.1 Detailed Description	137
7.21.2 Constructor & Destructor Documentation	137
7.21.2.1 NodeData()	137
7.21.3 Member Data Documentation	137
7.21.3.1 blengths	138
7.21.3.2 duplication	138
7.21.3.3 states	138
7.22 PhyloCounterData Class Reference	138
7.22.1 Detailed Description	139
7.22.2 Constructor & Destructor Documentation	139
7.22.2.1 PhyloCounterData()	139
7.22.3 Member Function Documentation	139
7.22.3.1 at()	139
7.22.3.2 begin()	139

7.22.3.3 empty()	. 139
7.22.3.4 end()	. 140
7.22.3.5 get_counters()	. 140
7.22.3.6 operator()()	. 140
7.22.3.7 push_back()	. 140
7.22.3.8 reserve()	. 140
7.22.3.9 shrink_to_fit()	. 140
7.22.3.10 size()	. 141
7.23 PhyloRuleDynData Class Reference	. 141
7.23.1 Detailed Description	. 141
7.23.2 Constructor & Destructor Documentation	. 141
7.23.2.1 PhyloRuleDynData()	. 141
$7.23.2.2 \sim PhyloRuleDynData() \ . \ . \ . \ . \ . \ . \ . \ . \ . \ $. 142
7.23.3 Member Data Documentation	. 142
7.23.3.1 counts	. 142
7.23.3.2 duplication	. 142
7.23.3.3 lb	. 142
7.23.3.4 pos	. 142
7.23.3.5 ub	. 142
7.24 PowerSet< Array_Type, Data_Rule_Type > Class Template Reference	. 143
7.24.1 Detailed Description	. 144
7.24.2 Constructor & Destructor Documentation	. 144
7.24.2.1 PowerSet() [1/3]	. 144
7.24.2.2 PowerSet() [2/3]	. 144
7.24.2.3 PowerSet() [3/3]	. 145
7.24.2.4 ~PowerSet()	. 145
7.24.3 Member Function Documentation	. 145
7.24.3.1 add_rule() [1/3]	. 145
7.24.3.2 add_rule() [2/3]	. 145
7.24.3.3 add_rule() [3/3]	. 145
7.24.3.4 begin()	. 146
7.24.3.5 calc()	. 146
7.24.3.6 end()	. 146
7.24.3.7 get_data()	. 146
7.24.3.8 get_data_ptr()	. 146
7.24.3.9 init_support()	. 147
7.24.3.10 operator[]()	. 147
7.24.3.11 reset()	. 147
7.24.3.12 size()	. 147
7.24.4 Member Data Documentation	. 147
7.24.4.1 coordinates_free	. 147
7.24.4.2 coordinates_locked	. 148

7.24.4.3 data	 148
7.24.4.4 EmptyArray	 148
7.24.4.5 M	 148
7.24.4.6 N	 148
7.24.4.7 rules	 149
7.24.4.8 rules_deleted	 149
7.25 Rule < Array_Type, Data_Type > Class Template Reference	 149
7.25.1 Detailed Description	 150
7.25.2 Constructor & Destructor Documentation	 150
7.25.2.1 Rule() [1/2]	 150
7.25.2.2 Rule() [2/2]	 150
7.25.2.3 ~Rule()	 150
7.25.3 Member Function Documentation	 151
7.25.3.1 D()	 151
7.25.3.2 operator()()	 151
7.26 Rules < Array_Type, Data_Type > Class Template Reference	 151
7.26.1 Detailed Description	 152
7.26.2 Constructor & Destructor Documentation	 152
7.26.2.1 Rules() [1/2]	 152
7.26.2.2 Rules() [2/2]	 152
7.26.2.3 ~Rules()	 153
7.26.3 Member Function Documentation	 153
7.26.3.1 add_rule() [1/3]	 153
7.26.3.2 add_rule() [2/3]	 153
7.26.3.3 add_rule() [3/3]	 153
7.26.3.4 clear()	 153
7.26.3.5 get_seq()	 153
7.26.3.6 operator()()	 154
7.26.3.7 operator=()	 154
7.26.3.8 size()	 155
7.27 StatsCounter< Array_Type, Data_Type > Class Template Reference $\dots \dots \dots \dots$	 155
7.27.1 Detailed Description	 155
7.27.2 Constructor & Destructor Documentation	 156
7.27.2.1 StatsCounter() [1/2]	 156
7.27.2.2 StatsCounter() [2/2]	 156
7.27.2.3 ~StatsCounter()	 156
7.27.3 Member Function Documentation	 156
7.27.3.1 add_counter() [1/2]	 156
7.27.3.2 add_counter() [2/2]	 157
7.27.3.3 count_all()	 157
7.27.3.4 count_current()	 157
7.27.3.5 count_init()	 157

7.27.3.6 get_counters()	. 157
7.27.3.7 reset_array()	. 157
7.27.3.8 set_counters()	. 158
7.28 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class Template Reference	
7.28.1 Detailed Description	. 161
7.28.2 Constructor & Destructor Documentation	. 161
7.28.2.1 Support() [1/3]	. 161
7.28.2.2 Support() [2/3]	. 161
7.28.2.3 Support() [3/3]	. 162
7.28.2.4 ~Support()	. 162
7.28.3 Member Function Documentation	. 162
7.28.3.1 add_counter() [1/2]	. 162
7.28.3.2 add_counter() [2/2]	. 162
7.28.3.3 add_rule() [1/2]	. 162
7.28.3.4 add_rule() [2/2]	. 163
7.28.3.5 add_rule_dyn() [1/2]	. 163
7.28.3.6 add_rule_dyn() [2/2]	. 163
7.28.3.7 calc()	. 163
7.28.3.8 eval_rules_dyn()	. 164
7.28.3.9 get_counters()	. 164
7.28.3.10 get_counts()	. 164
7.28.3.11 get_counts_ptr()	. 164
7.28.3.12 get_current_stats()	. 164
7.28.3.13 get_data()	. 165
7.28.3.14 get_rules()	. 165
7.28.3.15 get_rules_dyn()	. 165
7.28.3.16 init_support()	. 165
7.28.3.17 print()	. 165
7.28.3.18 reset_array() [1/2]	. 166
7.28.3.19 reset_array() [2/2]	. 166
7.28.3.20 set_counters()	. 166
7.28.3.21 set_rules()	. 166
7.28.3.22 set_rules_dyn()	. 166
7.28.4 Member Data Documentation	. 166
7.28.4.1 change_stats	. 167
7.28.4.2 coordinates_free	. 167
7.28.4.3 coordinates_locked	. 167
7.28.4.4 current_stats	. 167
7.28.4.5 delete_counters	. 167
7.28.4.6 delete_rules	. 168
7.28.4.7 delete rules dyn	. 168

7.28.4.8 M	 168
7.28.4.9 max_num_elements	 168
7.28.4.10 N	 168
7.29 vecHasher $<$ T $>$ Struct Template Reference	 169
7.29.1 Detailed Description	 169
7.29.2 Member Function Documentation	 169
7.29.2.1 operator()()	 169
8 File Documentation	171
8.1 include/barry/barray-bones.hpp File Reference	 171
8.1.1 Macro Definition Documentation	 172
8.1.1.1 BARRAY_BONES_HPP	 172
8.2 include/barry/barray-iterator.hpp File Reference	 172
8.3 include/barry/barray-meat-operators.hpp File Reference	 173
8.3.1 Macro Definition Documentation	 174
8.3.1.1 BARRY_BARRAY_MEAT_OPERATORS_HPP	 174
8.3.1.2 COL	 174
8.3.1.3 ROW	 174
8.3.2 Function Documentation	 174
8.3.2.1 checkdim_()	 174
8.4 include/barry/barray-meat.hpp File Reference	 175
8.4.1 Macro Definition Documentation	 176
8.4.1.1 COL	 176
8.4.1.2 ROW	 176
8.5 include/barry/barraycell-bones.hpp File Reference	 176
8.6 include/barry/barraycell-meat.hpp File Reference	 177
8.7 include/barry/barraydense-bones.hpp File Reference	 178
8.8 include/barry/barraydense-meet.hpp File Reference	 179
8.8.1 Macro Definition Documentation	 180
8.8.1.1 BARRY_BARRAYDENSE_MEAT_HPP	 180
8.8.1.2 COL	 181
8.8.1.3 POS	 181
8.8.1.4 ROW	 181
8.8.1.5 ZERO_CELL	 181
8.9 include/barry/barraydensecell-bones.hpp File Reference	 181
8.10 include/barry/barraydensecell-meat.hpp File Reference	 182
8.10.1 Macro Definition Documentation	 183
8.10.1.1 BARRY_BARRAYDENSECELL_MEAT_HPP	 183
8.10.1.2 POS	 183
8.11 include/barry/barrayvector-bones.hpp File Reference	 183
8.12 include/barry/barrayvector-meat.hpp File Reference	 184
8.12.1 Macro Definition Documentation	 185

8.12.1.1 BARRY_BARRAYVECTOR_MEAT_HPP
8.13 include/barry/barry-configuration.hpp File Reference
8.13.1 Macro Definition Documentation
8.13.1.1 BARRY_CHECK_SUPPORT
8.13.1.2 BARRY_ISFINITE
8.13.1.3 BARRY_MAX_NUM_ELEMENTS
8.13.1.4 BARRY_SAFE_EXP
8.13.1.5 printf_barry
8.13.2 Typedef Documentation
8.13.2.1 Map
8.14 include/barry/barry.hpp File Reference
8.14.1 Macro Definition Documentation
8.14.1.1 BARRY_HPP
8.14.1.2 BARRY_VERSION
8.14.1.3 COUNTER_FUNCTION
8.14.1.4 COUNTER_LAMBDA
8.14.1.5 RULE_FUNCTION
8.14.1.6 RULE_LAMBDA
8.15 include/barry/cell-bones.hpp File Reference
8.16 include/barry/cell-meat.hpp File Reference
8.17 include/barry/col-bones.hpp File Reference
8.18 include/barry/counters-bones.hpp File Reference
8.19 include/barry/counters-meat.hpp File Reference
8.20 include/barry/counters/network.hpp File Reference
8.20.1 Macro Definition Documentation
8.20.1.1 NET_C_DATA_IDX
8.20.1.2 NET_C_DATA_NUM
8.20.1.3 NETWORK_COUNTER
8.20.1.4 NETWORK_COUNTER_LAMBDA
8.20.1.5 NETWORK_RULE
8.20.1.6 NETWORK_RULE_LAMBDA
8.20.2 Typedef Documentation
8.20.2.1 NetCounter
8.20.2.2 NetCounters
8.20.2.3 NetModel
8.20.2.4 NetRule
8.20.2.5 NetRules
8.20.2.6 NetStatsCounter
8.20.2.7 NetSupport
8.20.2.8 Network
8.20.3 Function Documentation
8.20.3.1 rules_zerodiag()

8.21 include/barry/counters/phylo.hpp File Reference	200
8.21.1 Macro Definition Documentation	202
8.21.1.1 PHYLO_CHECK_MISSING	202
8.21.1.2 PHYLO_COUNTER_LAMBDA	202
8.21.1.3 PHYLO_RULE_DYN_LAMBDA	203
8.21.2 Typedef Documentation	203
8.21.2.1 PhyloArray	203
8.21.2.2 PhyloCounter	203
8.21.2.3 PhyloCounters	203
8.21.2.4 PhyloModel	203
8.21.2.5 PhyloPowerSet	204
8.21.2.6 PhyloRule	204
8.21.2.7 PhyloRuleData	204
8.21.2.8 PhyloRuleDyn	204
8.21.2.9 PhyloRules	204
8.21.2.10 PhyloRulesDyn	204
8.21.2.11 PhyloStatsCounter	205
8.21.2.12 PhyloSupport	205
8.21.3 Function Documentation	205
8.21.3.1 get_last_name()	205
8.22 include/barry/model-bones.hpp File Reference	205
8.22.1 Function Documentation	206
8.22.1.1 keygen_default()	206
8.23 include/barry/model-meat.hpp File Reference	207
8.23.1 Macro Definition Documentation	207
8.23.1.1 MODEL_TEMPLATE	207
8.23.1.2 MODEL_TEMPLATE_ARGS	208
8.23.1.3 MODEL_TYPE	208
8.23.2 Function Documentation	208
8.23.2.1 likelihood_()	208
8.23.2.2 MODEL_TEMPLATE() [1/2]	208
8.23.2.3 MODEL_TEMPLATE() [2/2]	209
8.23.2.4 update_normalizing_constant()	209
8.24 include/barry/models/geese.hpp File Reference	209
8.25 include/barry/models/geese/flock-bones.hpp File Reference	210
8.26 include/barry/models/geese/flock-meet.hpp File Reference	210
8.27 include/barry/models/geese/geese-bones.hpp File Reference	211
8.27.1 Macro Definition Documentation	211
8.27.1.1 INITIALIZED	212
8.27.2 Function Documentation	212
8.27.2.1 keygen_full()	212
8.27.2.2 RULE_FUNCTION()	212

8.27.2.3 vec_diff()	212
8.27.2.4 vector_caster()	212
8.28 include/barry/models/geese/geese-meat-constructors.hpp File Reference	213
8.29 include/barry/models/geese/geese-meat-likelihood.hpp File Reference	213
8.30 include/barry/models/geese/geese-meat-likelihood_exhaust.hpp File Reference	214
8.31 include/barry/models/geese/geese-meat-predict.hpp File Reference	215
8.32 include/barry/models/geese/geese-meat-predict_exhaust.hpp File Reference	215
8.33 include/barry/models/geese/geese-meat-predict_sim.hpp File Reference	216
8.34 include/barry/models/geese/geese-meat-simulate.hpp File Reference	216
8.35 include/barry/models/geese/geese-meat.hpp File Reference	217
8.36 include/barry/models/geese/geese-node-bones.hpp File Reference	217
8.37 include/barry/powerset-bones.hpp File Reference	218
8.38 include/barry/powerset-meat.hpp File Reference	219
8.39 include/barry/rules-bones.hpp File Reference	221
8.39.1 Function Documentation	222
8.39.1.1 rule_fun_default()	222
8.40 include/barry/rules-meat.hpp File Reference	222
8.41 include/barry/statscounter-bones.hpp File Reference	223
8.42 include/barry/statscounter-meat.hpp File Reference	224
8.43 include/barry/statsdb.hpp File Reference	225
8.44 include/barry/support-bones.hpp File Reference	226
8.45 include/barry/support-meat.hpp File Reference	228
8.45.1 Macro Definition Documentation	229
8.45.1.1 BARRY_SUPPORT_MEAT_HPP	230
8.45.1.2 SUPPORT_TEMPLATE	230
8.45.1.3 SUPPORT_TEMPLATE_ARGS	230
8.45.1.4 SUPPORT_TYPE	230
8.45.2 Function Documentation	230
8.45.2.1 calc_backend()	231
8.45.2.2 for()	231
8.45.2.3 if() [1/3]	231
8.45.2.4 if() [2/3]	231
8.45.2.5 if() [3/3]	231
8.45.2.6 insert_cell()	231
8.45.2.7 rm_cell()	232
8.45.2.8 SUPPORT_TEMPLATE() [1/17]	232
8.45.2.9 SUPPORT_TEMPLATE() [2/17]	232
8.45.2.10 SUPPORT_TEMPLATE() [3/17]	232
8.45.2.11 SUPPORT_TEMPLATE() [4/17]	232
8.45.2.12 SUPPORT_TEMPLATE() [5/17]	233
8.45.2.13 SUPPORT_TEMPLATE() [6/17]	233
8.45.2.14 SUPPORT_TEMPLATE() [7/17]	233

Index

8.45.2.15 SUPPORT_TEMPLATE	() [8/17]		 	 23
8.45.2.16 SUPPORT_TEMPLATE	() [9/17]		 	 23
8.45.2.17 SUPPORT_TEMPLATE	() [10/17]]	 	 23
8.45.2.18 SUPPORT_TEMPLATE	() [11/17]]	 	 234
8.45.2.19 SUPPORT_TEMPLATE	() [12/17]]	 	 234
8.45.2.20 SUPPORT_TEMPLATE	() [13/17]]	 	 23
8.45.2.21 SUPPORT_TEMPLATE	() [14/17]]	 	 234
8.45.2.22 SUPPORT_TEMPLATE	() [15/17]		 	 234
8.45.2.23 SUPPORT_TEMPLATE	() [16/17]]	 	 23
8.45.2.24 SUPPORT_TEMPLATE	() [17/17]		 	 23
8.45.3 Variable Documentation			 	 23
8.45.3.1 array_bank			 	 23
8.45.3.2 cfree			 	 23
8.45.3.3 counters			 	 23
8.45.3.4 counters			 	 230
8.45.3.5 delete_counters			 	 230
8.45.3.6 delete_rules			 	 230
8.45.3.7 delete_rules_dyn			 	 230
8.45.3.8 else			 	 230
8.45.3.9 f			 	 23
8.45.3.10 return			 	 23
8.45.3.11 rules			 	 23
8.45.3.12 rules			 	 23
8.45.3.13 rules_dyn			 	 23
8.45.3.14 stats_bank			 	 238
8.46 include/barry/typedefs.hpp File Reference			 	 238
8.46.1 Typedef Documentation			 	 240
8.46.1.1 Col_type			 	 240
8.46.1.2 Counter_fun_type			 	 240
8.46.1.3 Counts_type			 	 240
8.46.1.4 MapVec_type			 	 240
8.46.1.5 Row_type			 	 24
8.46.1.6 Rule_fun_type			 	 24
8.46.1.7 uint			 	 24
8.46.2 Function Documentation			 	 24
8.46.2.1 vec_equal()			 	 24
8.46.2.2 vec_equal_approx()			 	 24
8.46.2.3 vec_inner_prod()			 	 24
8.47 README.md File Reference			 	 24

243

Main Page

Barry: your to-go motif accountant

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

Examples

Counting statistics in a graph

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

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

2 Main Page

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

Efficient memory usage

. 1

2,] 1 .

Edges

C triads Mutuals

Transitive triads : 3
Isolates : 2

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

Code of Conduct

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

Module Index

2.1 Modules

Here is a list of all modules:

Counting			 																 		. (
Statistical Models																					
Network counters			 																 		. 10
Phylo counters			 																 		. 15
Phylo rules																					2-

4 Module Index

Class Index

3.1 Class List

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

BArray< Cell_Type, Data_Type >	
Baseline class for binary arrays	27
BArrayCell< Cell_Type, Data_Type >	41
BArrayCell_const< Cell_Type, Data_Type >	43
BArrayDense < Cell_Type, Data_Type >	
Baseline class for binary arrays	46
BArrayDenseCell< Cell_Type, Data_Type >	61
BArrayDenseCell_const< Cell_Type, Data_Type >	64
BArrayVector< Cell_Type, Data_Type >	
Row or column of a BArray	67
BArrayVector_const< Cell_Type, Data_Type >	71
Cell< Cell_Type >	
Entries in BArray. For now, it only has two members:	74
ConstBArrayRowlter< Cell_Type, Data_Type >	79
Counter< Array_Type, Data_Type >	
A counter function based on change statistics	81
Counters< Array_Type, Data_Type >	
Vector of counters	86
Entries < Cell_Type >	
A wrapper class to store source, target, val from a BArray object	90
Flock	
A Flock is a group of Geese	92
FreqTable < T >	
Database of statistics	99
Geese	
Annotated Phylo Model	101
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:	113
NetCounterData	
Data class used to store arbitrary uint or double vectors	127
NetworkData	
Data class for Networks	128
Node	
A single node for the model	131

6 Class Index

NodeData	
Data definition for the PhyloArray class	137
PhyloCounterData	138
PhyloRuleDynData	141
PowerSet< Array_Type, Data_Rule_Type >	
Powerset of a binary array	143
Rule < Array_Type, Data_Type >	
Rule for determining if a cell should be included in a sequence	149
Rules < Array_Type, Data_Type >	
Vector of objects of class Rule	151
StatsCounter< Array_Type, Data_Type >	
Count stats for a single Array	155
Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >	
Compute the support of sufficient statistics	158
vecHasher <t></t>	169

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

8 File Index

include/barry/models/geese/flock-meet.hpp	210
include/barry/models/geese/geese-bones.hpp	211
include/barry/models/geese/geese-meat-constructors.hpp	213
include/barry/models/geese/geese-meat-likelihood.hpp	213
include/barry/models/geese/geese-meat-likelihood_exhaust.hpp	214
include/barry/models/geese/geese-meat-predict.hpp	215
include/barry/models/geese/geese-meat-predict_exhaust.hpp	215
include/barry/models/geese/geese-meat-predict_sim.hpp	216
include/barry/models/geese/geese-meat-simulate.hpp	216
include/barry/models/geese/geese-meat.hpp	217
include/barry/models/geese/geese-node-bones hop	

Module Documentation

5.1 Counting

Classes

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

Data definition for the PhyloArray class.

class Counter< Array_Type, Data_Type >

A counter function based on change statistics.

5.1.1 Detailed Description

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

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

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

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

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

5.2 Statistical Models

Statistical models available in barry.

10 Module Documentation

Classes

• class Model < Array Type, Data Counter Type, Data Rule Type, Data Rule Dyn Type >

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

· class Flock

A Flock is a group of Geese.

class Geese

Annotated Phylo Model.

5.2.1 Detailed Description

Statistical models available in barry.

5.3 Network counters

Counters for network models.

Functions

void counter edges (NetCounters *counters)

Number of edges.

void counter isolates (NetCounters *counters)

Number of isolated vertices.

void counter_mutual (NetCounters *counters)

Number of mutual ties.

- void counter_istar2 (NetCounters *counters)
- void counter_ostar2 (NetCounters *counters)
- void counter_ttriads (NetCounters *counters)
- void counter_ctriads (NetCounters *counters)
- void counter_density (NetCounters *counters)
- void counter_idegree15 (NetCounters *counters)
- void counter_odegree15 (NetCounters *counters)
- void counter_absdiff (NetCounters *counters, uint attr_id, double alpha=1.0)

Sum of absolute attribute difference between ego and alter.

void counter_diff (NetCounters *counters, uint attr_id, double alpha=1.0, double tail_head=true)

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

- NETWORK_COUNTER (init_single_attr)
- void counter_nodeicov (NetCounters *counters, uint attr_id)
- void counter nodeocov (NetCounters *counters, uint attr id)
- void counter_nodecov (NetCounters *counters, uint attr_id)
- void counter nodematch (NetCounters *counters, uint attr id)
- void counter_idegree (NetCounters *counters, std::vector< uint > d)

Counts number of vertices with a given in-degree.

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

Counts number of vertices with a given out-degree.

void counter_degree (NetCounters *counters, std::vector< uint > d)

Counts number of vertices with a given out-degree.

void counter_css_partially_false_recip (NetCounters *counters, uint netsize, const std::vector< uint > &end)

Counts errors of commission.

5.3 Network counters

5.3.1 Detailed Description

Counters for network models.

Parameters

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

5.3.2 Function Documentation

5.3.2.1 counter_absdiff()

Sum of absolute attribute difference between ego and alter.

Definition at line 480 of file network.hpp.

5.3.2.2 counter_css_partially_false_recip()

Counts errors of commission.

< Start

 $< \mathsf{End}$

Definition at line 821 of file network.hpp.

5.3.2.3 counter_ctriads()

Definition at line 365 of file network.hpp.

12 Module Documentation

5.3.2.4 counter_degree()

Counts number of vertices with a given out-degree.

Definition at line 777 of file network.hpp.

5.3.2.5 counter_density()

Definition at line 413 of file network.hpp.

5.3.2.6 counter_diff()

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

Definition at line 524 of file network.hpp.

5.3.2.7 counter_edges()

Number of edges.

Definition at line 128 of file network.hpp.

5.3 Network counters

5.3.2.8 counter_idegree()

Counts number of vertices with a given in-degree.

Definition at line 680 of file network.hpp.

5.3.2.9 counter_idegree15()

Definition at line 434 of file network.hpp.

5.3.2.10 counter_isolates()

Number of isolated vertices.

Definition at line 149 of file network.hpp.

5.3.2.11 counter_istar2()

Definition at line 234 of file network.hpp.

5.3.2.12 counter_mutual()

Number of mutual ties.

Definition at line 187 of file network.hpp.

14 Module Documentation

5.3.2.13 counter_nodecov()

Definition at line 630 of file network.hpp.

5.3.2.14 counter_nodeicov()

Definition at line 586 of file network.hpp.

5.3.2.15 counter_nodematch()

Definition at line 652 of file network.hpp.

5.3.2.16 counter_nodeocov()

Definition at line 608 of file network.hpp.

5.3.2.17 counter_odegree()

Counts number of vertices with a given out-degree.

Definition at line 728 of file network.hpp.

5.4 Phylo counters 15

5.3.2.18 counter_odegree15()

Definition at line 456 of file network.hpp.

5.3.2.19 counter_ostar2()

Definition at line 255 of file network.hpp.

5.3.2.20 counter_ttriads()

Definition at line 279 of file network.hpp.

5.3.2.21 NETWORK_COUNTER()

Definition at line 568 of file network.hpp.

5.4 Phylo counters

Counters for phylogenetic modeling.

16 Module Documentation

Functions

void counter_overall_gains (PhyloCounters *counters, bool duplication=true)

Overall functional gains.

void counter_gains (PhyloCounters *counters, std::vector< uint > nfun, bool duplication=true)

Functional gains for a specific function (nfun).

void counter_gains_k_offspring (PhyloCounters *counters, std::vector< uint > nfun, uint k=1u, bool duplication=true)

k genes gain function nfun

void counter genes changing (PhyloCounters *counters, bool duplication=true)

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

• void counter_prop_genes_changing (PhyloCounters *counters, bool duplication=true)

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

• void counter_overall_loss (PhyloCounters *counters, bool duplication=true)

Overall functional loss.

void counter_maxfuns (PhyloCounters *counters, uint lb, uint ub, bool duplication=true)

Cap the number of functions per gene.

void counter_loss (PhyloCounters *counters, std::vector< uint > nfun, bool duplication=true)

Total count of losses for an specific function.

• void counter_overall_changes (PhyloCounters *counters, bool duplication=true)

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

• void counter_subfun (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)

Total count of Sub-functionalization events.

void counter_cogain (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)

Co-evolution (joint gain or loss)

void counter_longest (PhyloCounters *counters)

Longest branch mutates (either by gain or by loss)

• void counter_neofun (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)

Total number of neofunctionalization events.

• void counter neofun a2b (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)

Total number of neofunctionalization events.

void counter_co_opt (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)

Function co-opting.

5.4.1 Detailed Description

Counters for phylogenetic modeling.

Parameters

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

5.4.2 Function Documentation

5.4 Phylo counters

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 1184 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 814 of file phylo.hpp.

5.4.2.3 counter_gains()

Functional gains for a specific function (nfun).

Definition at line 179 of file phylo.hpp.

18 Module Documentation

5.4.2.4 counter_gains_k_offspring()

k genes gain function nfun

Definition at line 222 of file phylo.hpp.

5.4.2.5 counter_genes_changing()

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

Definition at line 296 of file phylo.hpp.

5.4.2.6 counter_longest()

Longest branch mutates (either by gain or by loss)

Definition at line 873 of file phylo.hpp.

5.4.2.7 counter_loss()

Total count of losses for an specific function.

Definition at line 612 of file phylo.hpp.

5.4 Phylo counters

5.4.2.8 counter_maxfuns()

Cap the number of functions per gene.

Definition at line 528 of file phylo.hpp.

5.4.2.9 counter_neofun()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 978 of file phylo.hpp.

5.4.2.10 counter_neofun_a2b()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1063 of file phylo.hpp.

5.4.2.11 counter_overall_changes()

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

Definition at line 661 of file phylo.hpp.

20 Module Documentation

5.4.2.12 counter_overall_gains()

Overall functional gains.

Total number of gains (irrespective of the function).

Definition at line 139 of file phylo.hpp.

5.4.2.13 counter_overall_loss()

Overall functional loss.

Definition at line 482 of file phylo.hpp.

5.4.2.14 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 367 of file phylo.hpp.

5.4.2.15 counter_subfun()

Total count of Sub-functionalization events.

It requires to specify data = {funA, funB}

Definition at line 728 of file phylo.hpp.

5.5 Phylo rules 21

5.5 Phylo rules

Rules for phylogenetic modeling.

Classes

• class PhyloRuleDynData

Functions

• void rule_dyn_limit_changes (PhyloSupport *support, uint pos, uint lb, uint ub, bool duplication=true) Overall functional gains.

5.5.1 Detailed Description

Rules for phylogenetic modeling.

Parameters

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

5.5.2 Function Documentation

5.5.2.1 rule_dyn_limit_changes()

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

Overall functional gains.

Parameters

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

22 Module Documentation

Returns

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

Definition at line 1317 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

- network
- phylo

6.2.1 Detailed Description

Tree class and Treelterator class.

6.3 barry::counters::network Namespace Reference

6.4 barry::counters::phylo Namespace Reference

6.5 CHECK Namespace Reference

Integer constants used to specify which cell should be check.

Variables

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

6.5.1 Detailed Description

Integer constants used to specify which cell should be check.

6.5.2 Variable Documentation

6.5.2.1 BOTH

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

Definition at line 20 of file typedefs.hpp.

6.5.2.2 NONE

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

Definition at line 21 of file typedefs.hpp.

6.5.2.3 ONE

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

Definition at line 22 of file typedefs.hpp.

6.5.2.4 TWO

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

Definition at line 23 of file typedefs.hpp.

6.6 EXISTS Namespace Reference

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

Variables

```
• const int BOTH = -1
```

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

6.6.1 Detailed Description

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

6.6.2 Variable Documentation

6.6.2.1 AS_ONE

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

Definition at line 38 of file typedefs.hpp.

6.6.2.2 AS_ZERO

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

Definition at line 37 of file typedefs.hpp.

6.6.2.3 BOTH

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

Definition at line 31 of file typedefs.hpp.

6.6.2.4 NONE

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

Definition at line 32 of file typedefs.hpp.

6.6.2.5 ONE

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

Definition at line 33 of file typedefs.hpp.

6.6.2.6 TWO

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

Definition at line 34 of file typedefs.hpp.

6.6.2.7 UKNOWN

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

Definition at line 36 of file typedefs.hpp.

Chapter 7

Class Documentation

7.1 BArray < Cell_Type, Data_Type > Class Template Reference

Baseline class for binary arrays.

#include <barray-bones.hpp>

Public Member Functions

- bool operator== (const BArray< Cell_Type, Data_Type > &Array_)
- ∼BArray ()
- void out_of_range (uint i, uint j) const
- Cell_Type get_cell (uint i, uint j, bool check_bounds=true) const
- 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

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.

• BArray ()

Zero-size array.

BArray (uint N_, uint M_)

Empty array.

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

Edgelist with data.

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

Edgelist with no data (simpler)

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

Move operator.

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

Set the data object.

- Data_Type * D ()
- const Data_Type * D () const
- · 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.

• BArray< Cell_Type, Data_Type > & operator+= (const std::pair< uint, uint > &coords)

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

Column/row wise interchange

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

Arithmetic operators

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

Public Attributes

bool visited = false

Friends

- class BArrayCell
 Cell_Type, Data_Type
- class BArrayCell_const< Cell_Type, Data_Type >

7.1.1 Detailed Description

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

Baseline class for binary arrays.

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

Template Parameters

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

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

7.1.2 Constructor & Destructor Documentation

7.1.2.1 BArray() [1/6]

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

Zero-size array.

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

7.1.2.2 BArray() [2/6]

Empty array.

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

7.1.2.3 BArray() [3/6]

Edgelist with data.

7.1.2.4 BArray() [4/6]

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

Edgelist with no data (simpler)

7.1.2.5 BArray() [5/6]

Copy constructor.

7.1.2.6 BArray() [6/6]

Move operator.

7.1.2.7 \sim BArray()

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

7.1.3 Member Function Documentation

7.1.3.1 clear()

7.1.3.2 col()

7.1.3.3 D() [1/2]

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

7.1.3.4 D() [2/2]

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

7.1.3.5 default_val()

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

7.1.3.6 flush_data()

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

7.1.3.7 get_cell()

7.1.3.8 get_col_vec() [1/2]

7.1.3.9 get_col_vec() [2/2]

7.1.3.10 get entries()

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

Get the edgelist.

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

Returns

Entries<Cell_Type>

7.1.3.11 get_row_vec() [1/2]

7.1.3.12 get_row_vec() [2/2]

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

7.1.3.13 insert_cell() [1/3]

7.1.3.14 insert_cell() [2/3]

7.1.3.15 insert_cell() [3/3]

7.1.3.16 is_empty()

7.1.3.17 ncol()

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

7.1.3.18 nnozero()

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

7.1.3.19 nrow()

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

7.1.3.20 operator()() [1/2]

7.1.3.21 operator()() [2/2]

7.1.3.22 operator*=()

7.1.3.23 operator+=() [1/3]

7.1.3.24 operator+=() [2/3]

7.1.3.25 operator+=() [3/3]

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/=()

7.1.3.30 operator=() [1/2]

Move assignment.

7.1.3.31 operator=() [2/2]

Assignment constructor.

7.1.3.32 operator==()

7.1.3.33 out_of_range()

7.1.3.34 print()

7.1.3.35 reserve()

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

7.1.3.36 resize()

7.1.3.37 rm_cell()

7.1.3.38 row()

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

7.1.3.39 set_data()

Set the data object.

Parameters

```
data_
delete_←
data_
```

7.1.3.40 swap_cells()

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

```
uint i0,
uint j0,
uint i1,
uint j1,
bool check_bounds = true,
int check_exists = CHECK::BOTH,
int * report = nullptr )
```

7.1.3.41 swap_cols()

7.1.3.42 swap_rows()

7.1.3.43 toggle_cell()

7.1.3.44 toggle_lock()

7.1.3.45 transpose()

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

7.1.3.46 zero col()

7.1.3.47 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 45 of file barray-bones.hpp.

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

• include/barry/barray-bones.hpp

7.2 BArrayCell< Cell Type, Data Type > Class Template Reference

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

Public Member Functions

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

7.2.1 Detailed Description

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

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

7.2.2 Constructor & Destructor Documentation

7.2.2.1 BArrayCell()

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

7.2.2.2 ∼BArrayCell()

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

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

7.2.3 Member Function Documentation

7.2.3.1 operator Cell_Type()

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

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

7.2.3.2 operator*=()

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

7.2.3.3 operator+=()

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

7.2.3.4 operator-=()

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

7.2.3.5 operator/=()

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

7.2.3.6 operator=()

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

7.2.3.7 operator==()

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

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

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

7.3 BArrayCell_const< Cell_Type, Data_Type > Class Template Reference

#include <barraycell-bones.hpp>

Public Member Functions

```
BArrayCell_const (const BArray < Cell_Type, Data_Type > *Array_, uint i_, uint j_, bool check_bounds=true)
~BArrayCell_const ()
operator Cell_Type () const
bool operator== (const Cell_Type &val) const
bool operator!= (const Cell_Type &val) const
bool operator < (const Cell_Type &val) const</li>
bool operator > (const Cell_Type &val) const
bool operator <= (const Cell_Type &val) const</li>
bool operator >= (const Cell_Type &val) const
bool operator >= (const Cell_Type &val) const
```

7.3.1 Detailed Description

```
\label{template} $$ \ensuremath{\sf template}$$ < \ensuremath{\sf type}$ = bool, typename Data_Type = bool> $$ \ensuremath{\sf class}$  \ensuremath{\sf BArrayCell\_const}< \ensuremath{\sf Cell\_Type}, Data_Type> $$
```

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

7.3.2 Constructor & Destructor Documentation

7.3.2.1 BArrayCell_const()

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

7.3.2.2 \sim BArrayCell_const()

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

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

7.3.3 Member Function Documentation

7.3.3.1 operator Cell_Type()

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

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

7.3.3.2 operator"!=()

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

7.3.3.3 operator<()

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

7.3.3.4 operator<=()

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

7.3.3.5 operator==()

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

7.3.3.6 operator>()

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

7.3.3.7 operator>=()

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

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

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

7.4 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
- const Row_type< Cell_Type > & row (uint i, bool check_bounds=true) const
- const Col_type< Cell_Type > & col (uint i, bool check_bounds=true) const
- Entries < Cell_Type > get_entries () const

Get the edgelist.

- · void transpose ()
- void clear (bool hard=true)
- void resize (uint N_, uint M_)
- void reserve ()
- · void print () const

Constructors

Parameters

N_	Number of rows
M_	Number of columns
source	An unsigned vector ranging from 0 to N_
target	An unsigned int vector ranging from 0 to M_
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.

- $\bullet \ \, \mathsf{BArrayDense} \ (\mathsf{BArrayDense} < \mathsf{Cell_Type}, \mathsf{Data_Type} > \&\&x) \ \mathsf{noexcept} \\$
 - 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 ()
- 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)
- BArrayCell< Cell_Type, Data_Type > operator() (uint i, uint j, bool check_bounds=true)
 const BArrayCell_const< Cell_Type, Data_Type > operator() (uint i, uint j, bool check_bounds=true) const
- void rm_cell (uint i, uint j, bool check_bounds=true, bool check_exists=true)
- void insert cell (uint i, uint i, const Cell < Cell Type > &v, bool check bounds, bool check exists)
- void insert_cell (uint i, uint i, 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 j1, uint j1, bool check bounds=true, int check exists=CHECK::BOTH, int *report=nullptr)
- void toggle cell (uint i, uint i, 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 BArrayCell
 Cell Type, Data Type
- class BArrayCell_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 arrays in which non-empty cells hold data of type Cell Type. The non-empty cells are stored by row and indexed using unordered_maps, i.e. std::vector< std↔ ::unordered_map<unsigned int,Cell_Type> >.

Template Parameters

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

Definition at line 22 of file 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 59 of file barraydense-bones.hpp.

7.4.2.2 BArrayDense() [2/6]

Empty array.

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

7.4.2.3 BArrayDense() [3/6]

Edgelist with data.

Definition at line 18 of file barraydense-meet.hpp.

7.4.2.4 BArrayDense() [4/6]

Edgelist with no data (simpler)

Definition at line 65 of file barraydense-meet.hpp.

7.4.2.5 BArrayDense() [5/6]

Copy constructor.

Definition at line 120 of file barraydense-meet.hpp.

7.4.2.6 BArrayDense() [6/6]

Move operator.

Definition at line 196 of file barraydense-meet.hpp.

7.4.2.7 ~BArrayDense()

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

Definition at line 273 of file barraydense-meet.hpp.

7.4.3 Member Function Documentation

7.4.3.1 clear()

Definition at line 1004 of file barraydense-meet.hpp.

7.4.3.2 col()

Definition at line 401 of file barraydense-meet.hpp.

7.4.3.3 D() [1/2]

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

Definition at line 297 of file barraydense-meet.hpp.

7.4.3.4 D() [2/2]

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

Definition at line 302 of file barraydense-meet.hpp.

7.4.3.5 default_val()

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

Definition at line 467 of file barraydense-meet.hpp.

7.4.3.6 get_cell()

Definition at line 317 of file barraydense-meet.hpp.

7.4.3.7 get_col_vec() [1/2]

Definition at line 377 of file barraydense-meet.hpp.

7.4.3.8 get_col_vec() [2/2]

Definition at line 362 of file barraydense-meet.hpp.

7.4.3.9 get_entries()

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

Get the edgelist.

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

Returns

```
Entries < Cell_Type >
```

Definition at line 412 of file barraydense-meet.hpp.

7.4.3.10 get_row_vec() [1/2]

Definition at line 346 of file barraydense-meet.hpp.

7.4.3.11 get_row_vec() [2/2]

Definition at line 331 of file barraydense-meet.hpp.

7.4.3.12 insert cell() [1/3]

```
template<typename Cell_Type , typename Data_Type >
void BArrayDense< Cell_Type, Data_Type >::insert_cell (
    uint i,
    uint j,
    Cell< Cell_Type > && v,
    bool check_bounds,
    bool check_exists ) [inline]
```

Definition at line 601 of file barraydense-meet.hpp.

7.4.3.13 insert_cell() [2/3]

```
template<typename Cell_Type , typename Data_Type >
void BArrayDense< Cell_Type, Data_Type >::insert_cell (
          uint i,
          uint j,
          Cell_Type v,
          bool check_bounds,
          bool check_exists ) [inline]
```

Definition at line 649 of file barraydense-meet.hpp.

7.4.3.14 insert_cell() [3/3]

```
template<typename Cell_Type , typename Data_Type >
void BArrayDense< Cell_Type, Data_Type >::insert_cell (
    uint i,
    uint j,
    const Cell< Cell_Type > & v,
    bool check_bounds,
    bool check_exists ) [inline]
```

Definition at line 553 of file barraydense-meet.hpp.

7.4.3.15 is_empty()

```
template<typename Cell_Type , typename Data_Type >
bool BArrayDense< Cell_Type, Data_Type >::is_empty (
          uint i,
          uint j,
          bool check_bounds = true ) const [inline]
```

Definition at line 432 of file barraydense-meet.hpp.

7.4.3.16 ncol()

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

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

7.4.3.17 nnozero()

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

Definition at line 462 of file barraydense-meet.hpp.

7.4.3.18 nrow()

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

Definition at line 452 of file barraydense-meet.hpp.

7.4.3.19 operator()() [1/2]

Definition at line 503 of file barraydense-meet.hpp.

7.4.3.20 operator()() [2/2]

Definition at line 512 of file barraydense-meet.hpp.

7.4.3.21 operator*=()

7.4.3.22 operator+=() [1/3]

7.4.3.23 operator+=() [2/3]

7.4.3.24 operator+=() [3/3]

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

7.4.3.25 operator-=() [1/3]

7.4.3.26 operator-=() [2/3]

7.4.3.27 operator-=() [3/3]

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

7.4.3.28 operator/=()

7.4.3.29 operator=() [1/2]

Move assignment.

Definition at line 212 of file barraydense-meet.hpp.

7.4.3.30 operator=() [2/2]

Assignment constructor.

Definition at line 156 of file barraydense-meet.hpp.

7.4.3.31 operator==()

Definition at line 254 of file barraydense-meet.hpp.

7.4.3.32 out_of_range()

```
template<typename Cell_Type , typename Data_Type >
void BArrayDense< Cell_Type, Data_Type >::out_of_range (
          uint i,
          uint j ) const [inline]
```

Definition at line 307 of file barraydense-meet.hpp.

7.4.3.33 print()

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

Definition at line 1072 of file barraydense-meet.hpp.

7.4.3.34 reserve()

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

Definition at line 1059 of file barraydense-meet.hpp.

7.4.3.35 resize()

Definition at line 1027 of file barraydense-meet.hpp.

7.4.3.36 rm_cell()

Definition at line 521 of file barraydense-meet.hpp.

7.4.3.37 row()

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

7.4.3.38 set_data()

Set the data object.

Parameters

data_	
delete_ <i>←</i>	
data_	

Definition at line 282 of file barraydense-meet.hpp.

7.4.3.39 swap_cells()

Definition at line 657 of file barraydense-meet.hpp.

7.4.3.40 swap_cols()

Definition at line 838 of file barraydense-meet.hpp.

7.4.3.41 swap_rows()

```
template<typename Cell_Type , typename Data_Type >
void BArrayDense< Cell_Type, Data_Type >::swap_rows (
          uint i0,
          uint i1,
          bool check_bounds = true ) [inline]
```

Definition at line 792 of file barraydense-meet.hpp.

7.4.3.42 toggle_cell()

```
template<typename Cell_Type , typename Data_Type >
void BArrayDense< Cell_Type, Data_Type >::toggle_cell (
          uint i,
          uint j,
          bool check_bounds = true,
          int check_exists = EXISTS::UKNOWN ) [inline]
```

Definition at line 758 of file barraydense-meet.hpp.

7.4.3.43 toggle_lock()

7.4.3.44 transpose()

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

Definition at line 947 of file barraydense-meet.hpp.

7.4.3.45 zero_col()

Definition at line 925 of file barraydense-meet.hpp.

7.4.3.46 zero_row()

Definition at line 906 of file barraydense-meet.hpp.

7.4.4 Friends And Related Function Documentation

7.4.4.1 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 barraydense-bones.hpp.

7.4.4.2 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 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 44 of file barraydense-bones.hpp.

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

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

7.5 BArrayDenseCell< Cell_Type, Data_Type > Class Template Reference

#include <barraydensecell-bones.hpp>

Public Member Functions

```
BArrayDenseCell (BArrayDense < Cell_Type, Data_Type > *Array_, uint i_, uint j_, bool check_bounds=true)
~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
```

7.5.1 Detailed Description

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

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

7.5.2 Constructor & Destructor Documentation

7.5.2.1 BArrayDenseCell()

Definition at line 16 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 28 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 44 of file barraydensecell-meat.hpp.

7.5.3.2 operator*=()

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

7.5.3.3 operator+=()

Definition at line 16 of file barraydensecell-meat.hpp.

7.5.3.4 operator-=()

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

7.5.3.5 operator/=()

Definition at line 37 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 49 of file barraydensecell-meat.hpp.

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

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

7.6 BArrayDenseCell_const< Cell_Type, Data_Type > Class Template Reference

#include <barraydensecell-bones.hpp>

Public Member Functions

- BArrayDenseCell_const (const BArrayDense < Cell_Type, Data_Type > *Array_, uint i_, uint j_, bool check
 _bounds=true)
- ∼BArrayDenseCell_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

7.6.1 Detailed Description

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

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

7.6.2 Constructor & Destructor Documentation

7.6.2.1 BArrayDenseCell_const()

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

7.6.2.2 ~BArrayDenseCell_const()

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

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

7.6.3 Member Function Documentation

7.6.3.1 operator Cell_Type()

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

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

7.6.3.2 operator"!=()

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

7.6.3.3 operator<()

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

7.6.3.4 operator<=()

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

7.6.3.5 operator==()

Definition at line 59 of file barraydensecell-meat.hpp.

7.6.3.6 operator>()

Definition at line 74 of file barraydensecell-meat.hpp.

7.6.3.7 operator>=()

Definition at line 84 of file barraydensecell-meat.hpp.

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

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

7.7 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

7.7.1 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.

7.7.2 Constructor & Destructor Documentation

7.7.2.1 BArrayVector()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
BArrayVector< Cell_Type, Data_Type >::BArrayVector (
```

```
BArray< Cell_Type, Data_Type > * Array_,
uint &dim_ uint & i_,
bool check_bounds = true ) [inline]
```

Construct a new BArrayVector object.

Parameters

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

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

7.7.2.2 ~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.

7.7.3 Member Function Documentation

7.7.3.1 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.

7.7.3.2 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.

7.7.3.3 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.

7.7.3.4 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.

7.7.3.5 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.

7.7.3.6 operator*=()

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

7.7.3.7 operator+=()

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

7.7.3.8 operator-=()

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

7.7.3.9 operator/=()

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

7.7.3.10 operator=()

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

7.7.3.11 operator==()

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

7.7.3.12 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

7.8 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

7.8.1 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.

7.8.2 Constructor & Destructor Documentation

7.8.2.1 BArrayVector_const()

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

7.8.2.2 ~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.

7.8.3 Member Function Documentation

7.8.3.1 begin()

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

7.8.3.2 end()

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

7.8.3.3 is col()

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

7.8.3.4 is_row()

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

7.8.3.5 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.

7.8.3.6 operator"!=()

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

7.8.3.7 operator<()

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

7.8.3.8 operator<=()

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

7.8.3.9 operator==()

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

7.8.3.10 operator>()

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

7.8.3.11 operator>=()

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

7.8.3.12 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

7.9 Cell < Cell Type > Class Template Reference

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

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

Public Member Functions

- Cell ()
- Cell (Cell_Type value_, bool visited_=false)
- ∼Cell ()
- Cell (const Cell
 Cell_Type > &arg)
- Cell< Cell_Type > & operator= (Cell< Cell_Type > &other)
- Cell (Cell< Cell_Type > &&arg) noexcept
- Cell< Cell_Type > & operator= (Cell< Cell_Type > &&other) noexcept
- void add (Cell_Type x)
- operator Cell_Type () const
- 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

7.9.1 Detailed Description

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

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

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

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

7.9.2 Constructor & Destructor Documentation

7.9.2.1 Cell() [1/7]

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

7.9.2.2 Cell() [2/7]

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

7.9.2.3 ∼CeII()

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

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

7.9.2.4 Cell() [3/7]

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

7.9.2.5 Cell() [4/7]

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

7.9.2.6 Cell() [5/7]

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

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

7.9.2.7 Cell() [6/7]

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

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

7.9.2.8 Cell() [7/7]

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

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

7.9.3 Member Function Documentation

7.9.3.1 add() [1/4]

7.9.3.2 add() [2/4]

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

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

7.9.3.3 add() [3/4]

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

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

7.9.3.4 add() [4/4]

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

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

7.9.3.5 operator Cell_Type()

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

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

7.9.3.6 operator"!=()

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

7.9.3.7 operator=() [1/2]

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

7.9.3.8 operator=() [2/2]

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

7.9.3.9 operator==()

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

7.9.4 Member Data Documentation

7.9.4.1 value

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

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

7.9.4.2 visited

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

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

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

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

7.10 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

7.10.1 Detailed Description

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

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

7.10.2 Constructor & Destructor Documentation

7.10.2.1 ConstBArrayRowlter()

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

7.10.2.2 ~ConstBArrayRowlter()

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

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

7.10.3 Member Data Documentation

7.10.3.1 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.

7.10.3.2 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.

7.10.3.3 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.

7.10.3.4 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

7.11 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)

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_=nullptr, Data_Type *data_=nullptr, bool delete_data_=false, std::string name_
 ="", std::string desc ="")
- Counter (const Counter < Array_Type, Data_Type > &counter_)
 Copy constructor.
- 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 = nullptr
    bool delete_data = false
    std::string name = ""
    std::string desc = ""
```

7.11.1 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.

7.11.2 Constructor & Destructor Documentation

7.11.2.1 Counter() [1/4]

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

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

7.11.2.2 Counter() [2/4]

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

7.11.2.3 Counter() [3/4]

Copy constructor.

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

7.11.2.4 Counter() [4/4]

Move constructor.

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

7.11.2.5 ~Counter()

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

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

7.11.3 Member Function Documentation

7.11.3.1 count()

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

7.11.3.2 init()

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

7.11.3.3 operator=() [1/2]

Copy assignment.

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

7.11.3.4 operator=() [2/2]

Move assignment.

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

7.11.4 Member Data Documentation

7.11.4.1 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.

7.11.4.2 data

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

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

7.11.4.3 delete_data

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

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

7.11.4.4 desc

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

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

7.11.4.5 init_fun

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

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

7.11.4.6 name

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

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

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

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

7.12 Counters < Array Type, Data Type > Class Template Reference

Vector of counters.

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

Public Member Functions

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

Copy constructor.

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

Move constructor.

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

Move assignment constructor.

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

Returns a pointer to a particular counter.

• std::size_t size () const noexcept

Number of counters in the set.

- void add_counter (Counter< Array_Type, Data_Type > &counter)
- void add_counter (Counter< Array_Type, Data_Type > *counter)
- void add_counter (Counter_fun_type< Array_Type, Data_Type > count_fun_, Counter_fun_type< Array_
 —
 Type, Data_Type > init_fun_=nullptr, Data_Type *data_=nullptr, bool delete_data_=false, std::string name
 —="", std::string desc_="")
- void clear ()

7.12.1 Detailed Description

Vector of counters.

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

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

7.12.2 Constructor & Destructor Documentation

7.12.2.1 Counters() [1/3]

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

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

7.12.2.2 ∼Counters()

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

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

7.12.2.3 Counters() [2/3]

Copy constructor.

Parameters



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

7.12.2.4 Counters() [3/3]

Move constructor.

Parameters



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

7.12.3 Member Function Documentation

7.12.3.1 add counter() [1/3]

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

7.12.3.2 add_counter() [2/3]

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

7.12.3.3 add_counter() [3/3]

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

7.12.3.4 clear()

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

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

7.12.3.5 operator=() [1/2]

Copy assignment constructor.

Parameters

counter←	
_	

Returns

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

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

7.12.3.6 operator=() [2/2]

Move assignment constructor.

Parameters



Returns

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

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

7.12.3.7 operator[]()

Returns a pointer to a particular counter.

Parameters

```
idx Id of the counter
```

Returns

Counter<Array_Type,Data_Type>*

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

7.12.3.8 size()

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

Number of counters in the set.

Returns

uint

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

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

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

7.13 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

7.13.1 Detailed Description

```
\label{template} \begin{tabular}{ll} template < typename Cell_Type > \\ class Entries < Cell_Type > \\ \end{tabular}
```

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

Template Parameters

```
Cell_Type Any type
```

Definition at line 59 of file typedefs.hpp.

7.13.2 Constructor & Destructor Documentation

7.13.2.1 Entries() [1/2]

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

Definition at line 65 of file typedefs.hpp.

7.13.2.2 Entries() [2/2]

Definition at line 66 of file typedefs.hpp.

7.13.2.3 ∼Entries()

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

Definition at line 73 of file typedefs.hpp.

7.13.3 Member Function Documentation

7.13.3.1 resize()

Definition at line 75 of file typedefs.hpp.

7.13.4 Member Data Documentation

7.13.4.1 source

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

Definition at line 61 of file typedefs.hpp.

7.13.4.2 target

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

Definition at line 62 of file typedefs.hpp.

7.13.4.3 val

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

Definition at line 63 of file typedefs.hpp.

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

• include/barry/typedefs.hpp

7.14 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 (bool verb=true)
- phylocounters::PhyloCounters * get_counters ()
- phylocounters::PhyloSupport * get_support ()
- 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) const noexcept
- void print () const

Public Attributes

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

7.14.1 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.

7.14.2 Constructor & Destructor Documentation

7.14.2.1 Flock()

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

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

7.14.2.2 ∼Flock()

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

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

7.14.3 Member Function Documentation

7.14.3.1 add_data()

Add a tree to the flock.

Parameters

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

Returns

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

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

7.14.3.2 colnames()

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

Definition at line 159 of file flock-meet.hpp.

7.14 Flock Class Reference 95

7.14.3.3 get_counters()

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

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

7.14.3.4 get_model()

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

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

7.14.3.5 get_support()

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

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

7.14.3.6 init()

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

7.14.3.7 likelihood_joint()

Returns the joint likelihood of the model.

Parameters

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

Returns

double

Definition at line 86 of file flock-meet.hpp.

7.14.3.8 nfuns()

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

Definition at line 111 of file flock-meet.hpp.

7.14.3.9 nleafs()

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

Definition at line 134 of file flock-meet.hpp.

7.14.3.10 nnodes()

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

Definition at line 123 of file flock-meet.hpp.

7.14.3.11 nterms()

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

Definition at line 146 of file flock-meet.hpp.

7.14.3.12 ntrees()

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

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

7.14.3.13 operator()()

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

Access the i-th geese element.

7.14 Flock Class Reference 97

Parameters

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

Returns

Geese *

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

7.14.3.14 parse_polytomies()

Definition at line 165 of file flock-meet.hpp.

7.14.3.15 print()

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

Definition at line 184 of file flock-meet.hpp.

7.14.3.16 set_seed()

Set the seed of the model.

Parameters

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

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

7.14.3.17 support_size()

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

Definition at line 153 of file flock-meet.hpp.

7.14.4 Member Data Documentation

7.14.4.1 dat

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

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

7.14.4.2 initialized

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

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

7.14.4.3 model

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

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

7.14.4.4 nfunctions

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

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

7.14.4.5 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-meet.hpp

7.15 FreqTable < T > Class Template Reference

Database of statistics.

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

Public Member Functions

- FreqTable ()
- ∼FreqTable ()
- void add (const std::vector< T > &x)
- Counts_type as_vector () const
- MapVec_type< T, uint > get_data () const
- const MapVec_type< T, uint > * get_data_ptr () const
- void clear ()
- void reserve (unsigned int n)
- void print () const
- size_t size () const noexcept

7.15.1 Detailed Description

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

Database of statistics.

This is mostly used in Support.

Definition at line 16 of file statsdb.hpp.

7.15.2 Constructor & Destructor Documentation

7.15.2.1 FreqTable()

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

Definition at line 28 of file statsdb.hpp.

7.15.2.2 ∼FreqTable()

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

Definition at line 29 of file statsdb.hpp.

7.15.3 Member Function Documentation

7.15.3.1 add()

Definition at line 47 of file statsdb.hpp.

7.15.3.2 as_vector()

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

Definition at line 61 of file statsdb.hpp.

7.15.3.3 clear()

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

Definition at line 83 of file statsdb.hpp.

7.15.3.4 get_data()

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

Definition at line 73 of file statsdb.hpp.

7.15.3.5 get_data_ptr()

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

Definition at line 78 of file statsdb.hpp.

7.16 Geese Class Reference 101

7.15.3.6 print()

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

Definition at line 102 of file statsdb.hpp.

7.15.3.7 reserve()

```
template<typename T >
void FreqTable< T >::reserve (
         unsigned int n ) [inline]
```

Definition at line 89 of file statsdb.hpp.

7.15.3.8 size()

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

Definition at line 126 of file statsdb.hpp.

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

• include/barry/statsdb.hpp

7.16 Geese Class Reference

Annotated Phylo Model.

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

Public Member Functions

- ∼Geese ()
- void init (bool verb=true)
- 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	geneid Id of the gene. It should be of length N.	
parent	Id of the parent gene. Also of length N	

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

Information about the model

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) const noexcept

Check polytomies and return the largest.

Geese prediction

Calculate the conditional probability

Parameters

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

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

Returns

std::vector< double > Returns the posterior probability

- std::vector< std::vector< double >> predict (const std::vector< double > &par, std::vector< std::vector< double >> *res_prob=nullptr, bool leave_one_out=false, bool only_annotated=false, bool use_reduced = sequence=true)
- std::vector < std::vector < double > > predict_backend (const std::vector < double > &par, bool use_← reduced sequence, const std::vector < 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() returns the computed support of the model.
```

- std::mt19937 * get_rengine ()
- phylocounters::PhyloCounters * get_counters ()
- phylocounters::PhyloModel * get model ()
- phylocounters::PhyloSupport * get_support ()

Public Attributes

- · unsigned int nfunctions
- std::map< unsigned int, Node > nodes
- barry::MapVec_type< unsigned int > map_to_nodes
- std::vector< unsigned int > sequence
- std::vector< unsigned int > reduced sequence
- bool initialized = false
- bool delete_rengine = false
- bool delete_support = false

7.16.1 Detailed Description

Annotated Phylo Model.

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

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

7.16.2 Constructor & Destructor Documentation

7.16.2.1 Geese() [1/4]

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

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

7.16.2.2 Geese() [2/4]

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

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

7.16.2.3 Geese() [3/4]

Definition at line 207 of file geese-meat-constructors.hpp.

7.16.2.4 Geese() [4/4]

Definition at line 280 of file geese-meat-constructors.hpp.

7.16.2.5 ∼Geese()

```
Geese::∼Geese ( ) [inline]
```

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

7.16.3 Member Function Documentation

7.16.3.1 calc_reduced_sequence()

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

Definition at line 274 of file geese-meat.hpp.

7.16.3.2 calc_sequence()

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

Definition at line 235 of file geese-meat.hpp.

7.16.3.3 colnames()

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

Names of the terms in the model.

Definition at line 375 of file geese-meat.hpp.

7.16.3.4 get_annotated_nodes()

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

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

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

7.16.3.5 get_counters()

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

Definition at line 539 of file geese-meat.hpp.

7.16.3.6 get_model()

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

Definition at line 544 of file geese-meat.hpp.

7.16.3.7 get_probabilities()

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

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

7.16.3.8 get_rengine()

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

Definition at line 534 of file geese-meat.hpp.

7.16 Geese Class Reference 107

7.16.3.9 get_states()

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

Powerset of a gene's possible states.

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

Returns

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

Definition at line 552 of file geese-meat.hpp.

7.16.3.10 get_support()

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

Definition at line 548 of file geese-meat.hpp.

7.16.3.11 inherit_support()

Definition at line 183 of file geese-meat.hpp.

7.16.3.12 init()

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

7.16.3.13 init_node()

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

7.16.3.14 likelihood()

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

7.16.3.15 likelihood_exhaust()

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

7.16.3.16 nannotations()

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

Number of annotations.

Definition at line 369 of file geese-meat.hpp.

7.16.3.17 nfuns()

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

Number of functions analyzed.

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

7.16.3.18 nleafs()

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

Number of leaf.

Definition at line 341 of file geese-meat.hpp.

7.16.3.19 nnodes()

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

Number of nodes (interior + leaf)

Definition at line 337 of file geese-meat.hpp.

7.16.3.20 nterms()

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

Number of terms included.

Definition at line 351 of file geese-meat.hpp.

7.16.3.21 observed_counts()

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

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

7.16.3.22 operator=() [1/2]

7.16.3.23 operator=() [2/2]

7.16.3.24 parse_polytomies()

Check polytomies and return the largest.

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

7.16.3.25 predict()

Definition at line 240 of file geese-meat-predict.hpp.

7.16.3.26 predict_backend()

< True if the array belongs to the set

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

7.16.3.27 predict_exhaust()

Definition at line 5 of file geese-meat-predict exhaust.hpp.

7.16.3.28 predict_exhaust_backend()

Definition at line 47 of file geese-meat-predict exhaust.hpp.

7.16.3.29 predict_sim()

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

7.16 Geese Class Reference 111

7.16.3.30 print()

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

Prints information about the GEESE.

Definition at line 516 of file geese-meat.hpp.

7.16.3.31 print_observed_counts()

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

Definition at line 454 of file geese-meat.hpp.

7.16.3.32 set_seed()

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

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

7.16.3.33 simulate()

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

7.16.3.34 support_size()

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

Number of unique sets of sufficient stats.

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

7.16.3.35 update_annotations()

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

Definition at line 206 of file geese-meat.hpp.

7.16.4 Member Data Documentation

7.16.4.1 delete_rengine

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

Definition at line 117 of file geese-bones.hpp.

7.16.4.2 delete_support

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

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

7.16.4.3 initialized

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

Definition at line 116 of file geese-bones.hpp.

7.16.4.4 map_to_nodes

```
\verb|barry::MapVec_type<| unsigned int > Geese::map_to_nodes|
```

Definition at line 109 of file geese-bones.hpp.

7.16.4.5 nfunctions

unsigned int Geese::nfunctions

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

7.16.4.6 nodes

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

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

7.16.4.7 reduced sequence

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

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

7.16.4.8 sequence

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

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

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

- include/barry/models/geese/geese-bones.hpp
- include/barry/models/geese/geese-meat-constructors.hpp
- include/barry/models/geese/geese-meat-likelihood.hpp
- include/barry/models/geese/geese-meat-likelihood_exhaust.hpp
- include/barry/models/geese/geese-meat-predict.hpp
- include/barry/models/geese/geese-meat-predict_exhaust.hpp
- $\bullet \ \ include/barry/models/geese/geese-meat-predict_sim.hpp$
- include/barry/models/geese/geese-meat-simulate.hpp
- include/barry/models/geese/geese-meat.hpp

7.17 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 ()

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

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

- void add_counter (Counter< Array_Type, Data_Counter_Type > &counter)
- void add counter (Counter< Array Type, Data Counter Type > *counter)
- void add_counter (Counter_fun_type< Array_Type, Data_Counter_Type > count_fun_, Counter_fun_type
 Array_Type, Data_Counter_Type > init_fun_=nullptr, Data_Counter_Type *data_=nullptr, bool delete_
 data =false)
- void set counters (Counters < Array Type, Data Counter Type > *counters)

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

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

- void add_rule (Rule < Array_Type, Data_Rule_Type > &rule)
- void add_rule (Rule < Array_Type, Data_Rule_Type > *rule)
- void add_rule (Rule_fun_type< Array_Type, Data_Rule_Type > count_fun_, Data_Rule_Type *data_
 =nullptr, bool delete_data_=false)
- void set_rules (Rules< Array_Type, Data_Rule_Type > *rules_)
- void add_rule_dyn (Rule< Array_Type, Data_Rule_Dyn_Type > &rule)
- void add rule dyn (Rule < Array Type, Data Rule Dyn Type > *rule)
- void 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 > ¶ms, const std::vector< 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< 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.

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

7.17.1 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 47 of file model-bones.hpp.

7.17.2 Constructor & Destructor Documentation

7.17.2.1 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 ( )
```

7.17.2.2 Model() [2/3]

```
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 (
    uint size_ )
```

7.17.2.3 Model() [3/3]

7.17.2.4 ∼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 149 of file model-bones.hpp.

7.17.3 Member Function Documentation

7.17.3.1 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.

7.17.3.2 add_counter() [1/3]

7.17.3.3 add_counter() [2/3]

7.17.3.4 add counter() [3/3]

7.17.3.5 add_rule() [1/3]

```
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 )
```

7.17.3.6 add rule() [2/3]

```
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 )
```

7.17.3.7 add_rule() [3/3]

7.17.3.8 add rule dyn() [1/3]

7.17.3.9 add_rule_dyn() [2/3]

7.17.3.10 add_rule_dyn() [3/3]

7.17.3.11 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
```

7.17.3.12 conditional prob()

Conditional probability ("Gibbs sampler")

Computes the conditional probability of observing $P\{Y(i,j) = | Y^{\wedge}C, \text{ theta}\}$, 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

7.17.3.13 gen_key()

7.17.3.14 get_counters()

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

7.17.3.15 get_norm_const()

7.17.3.16 get_pset()

7.17.3.17 get_pset_stats()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftrightarrow Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> const std::vector< std::vector< double > >* Model< Array_Type, Data_Counter_Type, Data_Rule\leftrightarrow _Type, Data_Rule_Dyn_Type >::get_pset_stats ( const uint & i)
```

7.17.3.18 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
```

7.17.3.19 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 ()
```

7.17.3.20 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 ()
```

7.17.3.21 get_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>* Model< Array_Type,
Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::get_support ( )
```

7.17.3.22 likelihood() [1/3]

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarrow Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> double Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::likelihood ( const std::vector< double > & params, const Array_Type & Array_, int i = -1, bool as_log = false)
```

7.17.3.23 likelihood() [2/3]

7.17.3.24 likelihood() [3/3]

7.17.3.25 likelihood_total()

7.17.3.26 nterms()

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

7.17.3.27 operator=()

7.17.3.28 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.

7.17.3.29 print_stats()

```
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_stats (
    uint i) const
```

7.17.3.30 sample() [1/2]

7.17.3.31 sample() [2/2]

7.17.3.32 set_counters()

7.17.3.33 set_keygen()

7.17.3.34 set_rengine()

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

Definition at line 119 of file model-bones.hpp.

7.17.3.35 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_ )
```

7.17.3.36 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_ )
```

7.17.3.37 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 129 of file model-bones.hpp.

7.17.3.38 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]
```

7.17.3.39 size_unique()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
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_unique ( ) const [noexcept]
```

7.17.3.40 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]
```

7.17.3.41 support_size()

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

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

• include/barry/model-bones.hpp

7.18 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

7.18.1 Detailed Description

Data class used to store arbitrary uint or double vectors.

Definition at line 61 of file network.hpp.

7.18.2 Constructor & Destructor Documentation

7.18.2.1 NetCounterData() [1/2]

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

Definition at line 67 of file network.hpp.

7.18.2.2 NetCounterData() [2/2]

Definition at line 68 of file network.hpp.

7.18.2.3 ∼NetCounterData()

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

Definition at line 73 of file network.hpp.

7.18.3 Member Data Documentation

7.18.3.1 indices

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

Definition at line 64 of file network.hpp.

7.18.3.2 numbers

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

Definition at line 65 of file network.hpp.

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

• include/barry/counters/network.hpp

7.19 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

7.19.1 Detailed Description

Data class for Networks.

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

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

Definition at line 24 of file network.hpp.

7.19.2 Constructor & Destructor Documentation

7.19.2.1 NetworkData() [1/3]

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

Definition at line 30 of file network.hpp.

7.19.2.2 NetworkData() [2/3]

Constructor using a single attribute.

Parameters

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

Definition at line 38 of file network.hpp.

7.19.2.3 NetworkData() [3/3]

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

Constructor using multiple attributes.

Parameters

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

Definition at line 50 of file network.hpp.

7.19.2.4 ~NetworkData()

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

Definition at line 56 of file network.hpp.

7.19.3 Member Data Documentation

7.19.3.1 directed

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

Definition at line 27 of file network.hpp.

7.20 Node Class Reference 131

7.19.3.2 vertex_attr

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

Definition at line 28 of file network.hpp.

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

• include/barry/counters/network.hpp

7.20 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.

7.20.1 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.

7.20.2 Constructor & Destructor Documentation

7.20.2.1 Node() [1/5]

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

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

7.20 Node Class Reference 133

7.20.2.2 Node() [2/5]

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

Definition at line 56 of file geese-node-bones.hpp.

7.20.2.3 Node() [3/5]

```
Node::Node (
          unsigned int id_,
          unsigned int ord_,
          std::vector< unsigned int > annotations_,
          bool duplication_ ) [inline]
```

Definition at line 62 of file geese-node-bones.hpp.

7.20.2.4 Node() [4/5]

Definition at line 69 of file geese-node-bones.hpp.

7.20.2.5 Node() [5/5]

Definition at line 83 of file geese-node-bones.hpp.

7.20.2.6 ∼Node()

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

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

7.20.3 Member Function Documentation

7.20.3.1 get_parent()

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

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

7.20.3.2 is_leaf()

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

Definition at line 109 of file geese-node-bones.hpp.

7.20.3.3 noffspring()

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

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

7.20.4 Member Data Documentation

7.20.4.1 annotations

std::vector< unsigned int > Node::annotations

Observed annotations (only defined for Geese)

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

7.20.4.2 array

phylocounters::PhyloArray Node::array

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

7.20 Node Class Reference 135

7.20.4.3 arrays

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

Arrays given all possible states.

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

7.20.4.4 duplication

```
bool Node::duplication
```

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

7.20.4.5 id

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

Id of the node (as specified in the input)

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

7.20.4.6 narray

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

ID of the array in the model.

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

7.20.4.7 offspring

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

Offspring nodes.

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

7.20.4.8 ord

unsigned int Node::ord

Order in which the node was created.

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

7.20.4.9 parent

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

Parent node.

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

7.20.4.10 probability

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

The probability of observing each state.

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

7.20.4.11 subtree_prob

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

Induced subtree probabilities.

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

7.20.4.12 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

7.21 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
```

7.21.1 Detailed Description

Data definition for the PhyloArray class.

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

This holds basic information about a given node.

Definition at line 23 of file phylo.hpp.

7.21.2 Constructor & Destructor Documentation

7.21.2.1 NodeData()

Definition at line 43 of file phylo.hpp.

7.21.3 Member Data Documentation

7.21.3.1 blengths

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

Branch length.

Definition at line 29 of file phylo.hpp.

7.21.3.2 duplication

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

Definition at line 39 of file phylo.hpp.

7.21.3.3 states

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

State of the parent node.

Definition at line 34 of file phylo.hpp.

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

• include/barry/counters/phylo.hpp

7.22 PhyloCounterData Class Reference

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

Public Member Functions

- PhyloCounterData (std::vector< uint > data_, std::vector< double > *counters_=nullptr)
- uint at (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 ()

7.22.1 Detailed Description

Definition at line 54 of file phylo.hpp.

7.22.2 Constructor & Destructor Documentation

7.22.2.1 PhyloCounterData()

Definition at line 60 of file phylo.hpp.

7.22.3 Member Function Documentation

7.22.3.1 at()

Definition at line 65 of file phylo.hpp.

7.22.3.2 begin()

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

Definition at line 72 of file phylo.hpp.

7.22.3.3 empty()

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

Definition at line 75 of file phylo.hpp.

7.22.3.4 end()

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

Definition at line 73 of file phylo.hpp.

7.22.3.5 get_counters()

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

Definition at line 76 of file phylo.hpp.

7.22.3.6 operator()()

Definition at line 66 of file phylo.hpp.

7.22.3.7 push_back()

Definition at line 68 of file phylo.hpp.

7.22.3.8 reserve()

Definition at line 67 of file phylo.hpp.

7.22.3.9 shrink_to_fit()

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

Definition at line 69 of file phylo.hpp.

7.22.3.10 size()

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

Definition at line 70 of file phylo.hpp.

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

• include/barry/counters/phylo.hpp

7.23 PhyloRuleDynData Class Reference

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

Public Member Functions

- PhyloRuleDynData (const std::vector< double > *counts_, uint pos_, uint lb_, uint ub_, bool duplication_)
- ∼PhyloRuleDynData ()

Public Attributes

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

7.23.1 Detailed Description

Definition at line 1289 of file phylo.hpp.

7.23.2 Constructor & Destructor Documentation

7.23.2.1 PhyloRuleDynData()

Definition at line 1296 of file phylo.hpp.

7.23.2.2 ~PhyloRuleDynData()

PhyloRuleDynData::~PhyloRuleDynData () [inline]

Definition at line 1305 of file phylo.hpp.

7.23.3 Member Data Documentation

7.23.3.1 counts

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

Definition at line 1291 of file phylo.hpp.

7.23.3.2 duplication

bool PhyloRuleDynData::duplication

Definition at line 1295 of file phylo.hpp.

7.23.3.3 lb

```
uint PhyloRuleDynData::lb
```

Definition at line 1293 of file phylo.hpp.

7.23.3.4 pos

```
uint PhyloRuleDynData::pos
```

Definition at line 1292 of file phylo.hpp.

7.23.3.5 ub

```
uint PhyloRuleDynData::ub
```

Definition at line 1294 of file phylo.hpp.

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

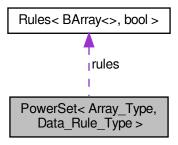
• include/barry/counters/phylo.hpp

7.24 PowerSet < Array_Type, Data_Rule_Type > Class Template Reference

Powerset of a binary array.

#include <powerset-bones.hpp>

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



Public Member Functions

- void init_support ()
- void calc ()
- void reset (uint N_, uint M_)

Construct and destroy a PowerSet object

- PowerSet ()
- PowerSet (uint N_, uint M_)
- PowerSet (const Array_Type & array)
- ∼PowerSet ()

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

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

- void add_rule (Rule < Array_Type, Data_Rule_Type > &rule)
- void add_rule (Rule < Array_Type, Data_Rule_Type > *rule)
- void add_rule (Rule_fun_type< Array_Type, Data_Rule_Type > count_fun_, Data_Rule_Type *data_← =nullptr, bool delete_data_=false)

Getter functions

- const std::vector< Array_Type > * get_data_ptr () const
- std::vector< Array_Type > get_data () const
 std::vector< Array_Type >::iterator begin ()
- std::vector< Array_Type >::iterator end ()
- std::size_t size () const noexcept
- const Array_Type & operator[] (const unsigned int &i) const

Public Attributes

- Array_Type EmptyArray
- std::vector< Array_Type > data
- Rules< Array_Type, Data_Rule_Type > * rules
- uint N
- uint M
- bool rules_deleted = false
- std::vector< std::pair< uint, uint >> coordinates_free
- std::vector< std::pair< uint, uint >> coordinates_locked

7.24.1 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.

7.24.2 Constructor & Destructor Documentation

7.24.2.1 PowerSet() [1/3]

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

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

7.24.2.2 PowerSet() [2/3]

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

7.24.2.3 PowerSet() [3/3]

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

7.24.2.4 ∼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.

7.24.3 Member Function Documentation

7.24.3.1 add_rule() [1/3]

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

7.24.3.2 add_rule() [2/3]

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

7.24.3.3 add_rule() [3/3]

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

7.24.3.4 begin()

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

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

7.24.3.5 calc()

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

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

7.24.3.6 end()

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

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

7.24.3.7 get_data()

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

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

7.24.3.8 get_data_ptr()

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

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

7.24.3.9 init_support()

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

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

7.24.3.10 operator[]()

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

7.24.3.11 reset()

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

7.24.3.12 size()

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

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

7.24.4 Member Data Documentation

7.24.4.1 coordinates free

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

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

7.24.4.2 coordinates_locked

template<typename Array_Type = BArray<>, typename Data_Rule_Type = bool>
std::vector< std::pair<uint,uint> > PowerSet< Array_Type, Data_Rule_Type >::coordinates_← locked

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

7.24.4.3 data

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

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

7.24.4.4 EmptyArray

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

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

7.24.4.5 M

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

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

7.24.4.6 N

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

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

7.24.4.7 rules

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

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

7.24.4.8 rules_deleted

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

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

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

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

7.25 Rule < Array_Type, Data_Type > Class Template Reference

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

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

Public Member Functions

- ∼Rule ()
- Data_Type * D ()

Read/Write access to the data.

• bool operator() (const Array_Type &a, uint i, uint j)

Construct a new Rule object

Construct a new Rule object

Parameters

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

- Rule ()
- Rule (Rule_fun_type< Array_Type, Data_Type > fun_, Data_Type *dat_=nullptr, bool delete_dat_=false)

7.25.1 Detailed Description

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

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

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

Template Parameters

Array_Type	An object of class BArray.
Data_Type	Any type.

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

7.25.2 Constructor & Destructor Documentation

7.25.2.1 Rule() [1/2]

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

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

7.25.2.2 Rule() [2/2]

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

7.25.2.3 ∼Rule()

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

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

7.25.3 Member Function Documentation

7.25.3.1 D()

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

Read/Write access to the data.

7.25.3.2 operator()()

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

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

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

7.26 Rules < Array_Type, Data_Type > Class Template Reference

Vector of objects of class Rule.

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

Public Member Functions

- Rules ()
- Rules (const Rules < Array_Type, Data_Type > &rules_)
- Rules< Array_Type, Data_Type > operator= (const Rules< Array_Type, Data_Type > &rules_)
- ∼Rules ()
- uint size () const noexcept
- bool operator() (const Array_Type &a, uint i, uint j)

Check whether a given cell is free or locked.

- void clear ()
- void get_seq (const Array_Type &a, std::vector< std::pair< uint, uint >> *free, std::vector< std::pair< uint, uint >> *locked=nullptr)

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

Rule adding

Parameters

```
rule
```

- void add rule (Rule < Array Type, Data Type > &rule)
- void add rule (Rule < Array Type, Data Type > *rule)
- void add_rule (Rule_fun_type< Array_Type, Data_Type > rule_, Data_Type *data_=nullptr, bool delete ← data = false)

7.26.1 Detailed Description

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

Vector of objects of class Rule.

Template Parameters

Array_Type	An object of class BArray
Data_Type	Any type.

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

7.26.2 Constructor & Destructor Documentation

7.26.2.1 Rules() [1/2]

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

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

7.26.2.2 Rules() [2/2]

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

7.26.2.3 ∼Rules()

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

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

7.26.3 Member Function Documentation

7.26.3.1 add_rule() [1/3]

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

7.26.3.2 add_rule() [2/3]

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

7.26.3.3 add_rule() [3/3]

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

7.26.3.4 clear()

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

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

7.26.3.5 get_seq()

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

Parameters

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

Returns

Nothing.

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

7.26.3.6 operator()()

Check whether a given cell is free or locked.

Parameters

а	A BArray object
i	row position
j	col position

Returns

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

Definition at line 111 of file rules-meat.hpp.

7.26.3.7 operator=()

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

7.26.3.8 size()

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

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

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

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

7.27 StatsCounter< Array_Type, Data_Type > Class Template Reference

Count stats for a single Array.

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

Public Member Functions

StatsCounter (const Array Type *Array)

Creator of a StatsCounter

• StatsCounter ()

Can be created without setting the array.

- \sim StatsCounter ()
- void reset_array (const Array_Type *Array_)

Changes the reference array for the counting.

- void add_counter (Counter < Array_Type, Data_Type > *f_)
- void add_counter (Counter< Array_Type, Data_Type > f_)
- void set_counters (Counters< Array_Type, Data_Type > *counters_)
- void count_init (uint i, uint j)

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

- void count_current (uint i, uint j)
- std::vector< double > count all ()
- Counters < Array_Type, Data_Type > * get_counters ()

7.27.1 Detailed Description

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

Count stats for a single Array.

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

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

7.27.2 Constructor & Destructor Documentation

7.27.2.1 StatsCounter() [1/2]

Creator of a StatsCounter

Parameters

Array←	A const pointer to a BArray.	
_		

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

7.27.2.2 StatsCounter() [2/2]

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

Can be created without setting the array.

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

7.27.2.3 ~StatsCounter()

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

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

7.27.3 Member Function Documentation

7.27.3.1 add_counter() [1/2]

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

7.27.3.2 add_counter() [2/2]

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

7.27.3.3 count all()

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

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

7.27.3.4 count_current()

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

7.27.3.5 count init()

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

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

7.27.3.6 get_counters()

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

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

7.27.3.7 reset_array()

Changes the reference array for the counting.

158 Class Documentation

Parameters

Array←	A pointer to an array of class Array_Type.

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

7.27.3.8 set_counters()

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

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

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

7.28 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class Template Reference

Compute the support of sufficient statistics.

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

Public Member Functions

Support (const Array_Type &Array_)

Constructor passing a reference Array.

Support (uint N_, uint M_)

Constructor specifying the dimensions of the array (empty).

- Support ()
- ∼Support ()
- void init_support (std::vector< Array_Type > *array_bank=nullptr, std::vector< std::vector< double > >
 *stats_bank=nullptr)
- void calc (std::vector< Array_Type > *array_bank=nullptr, std::vector< std::vector< double > > *stats_bank=nullptr, unsigned int max_num_elements_=0u)

Computes the entire support.

- Counts_type get_counts () const
- const MapVec_type * get_counts_ptr () const
- std::vector< double > * get_current_stats ()

List current statistics.

- void print () const
- const FreqTable & get_data () const
- Counters < Array_Type, Data_Counter_Type > * get_counters ()

Vector of couter functions.

• Rules< Array_Type, Data_Rule_Type > * get_rules ()

Vector of static rules (cells to iterate).

Rules< Array_Type, Data_Rule_Dyn_Type > * get_rules_dyn ()

Vector of dynamic rules (to include/exclude a realizaton).

Resets the support calculator

If needed, the counters of a support object can be reused.

160 Class Documentation

Parameters

Array←	New array over which the support will be computed.

- · void reset_array ()
- void reset_array (const Array_Type &Array_)

Manage counters

Parameters

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

- void add_counter (Counter< Array_Type, Data_Counter_Type > *f_)
- void add_counter (Counter< Array_Type, Data_Counter_Type > f_)
- void set_counters (Counters < Array_Type, Data_Counter_Type > *counters_)

Manage rules

Parameters

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

- void add_rule (Rule < Array_Type, Data_Rule_Type > *f_)
- void add_rule (Rule < Array_Type, Data_Rule_Type > f_)
- void set_rules (Rules< Array_Type, Data_Rule_Type > *rules_)
- void add_rule_dyn (Rule< Array_Type, Data_Rule_Dyn_Type > *f_)
- void add_rule_dyn (Rule < Array_Type, Data_Rule_Dyn_Type > f_)
- void set_rules_dyn (Rules < Array_Type, Data_Rule_Dyn_Type > *rules_)
- 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< std::pair< uint, uint >> coordinates_free
- $\bullet \ \ \mathsf{std} :: \mathsf{vector} < \mathsf{std} :: \mathsf{pair} < \mathsf{uint}, \ \mathsf{uint} > > \mathsf{coordinates_locked} \\$
- std::vector< std::vector< double >> change stats

7.28.1 Detailed Description

template < typename Array_Type = BArray <>>, typename Data_Counter_Type = bool, typename Data_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>

class Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >

Compute the support of sufficient statistics.

Given an array and a set of counters, this object iterates throughout the support set of the Array while at the same time computing the support of the sufficient statitics.

The members rule and rule_dyn allow constraining the support. The first will establish which cells of the array will be used to iterate, for example, in the case of social networks, self-loops are not allowed, so the entire diagonal would be fixed to zero, reducing the size of the support.

In the case of $rule_dyn$, the function will stablish dynamically whether the current state will be included in the counts or not. For example, this set of rules can be used to constrain the support to networks that have a prescribed degree sequence.

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

7.28.2 Constructor & Destructor Documentation

7.28.2.1 Support() [1/3]

Constructor passing a reference Array.

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

7.28.2.2 Support() [2/3]

Constructor specifying the dimensions of the array (empty).

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

162 Class Documentation

7.28.2.3 Support() [3/3]

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

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

7.28.2.4 ~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 92 of file support-bones.hpp.

7.28.3 Member Function Documentation

7.28.3.1 add counter() [1/2]

7.28.3.2 add_counter() [2/2]

7.28.3.3 add_rule() [1/2]

7.28.3.4 add_rule() [2/2]

7.28.3.5 add_rule_dyn() [1/2]

7.28.3.6 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_- )
```

7.28.3.7 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< 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.

164 Class Documentation

7.28.3.8 eval_rules_dyn()

7.28.3.9 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.

7.28.3.10 get_counts()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Counts_type Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >←
::get_counts () const
```

7.28.3.11 get_counts_ptr()

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

7.28.3.12 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.

7.28.3.13 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
```

7.28.3.14 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).

7.28.3.15 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).

7.28.3.16 init support()

7.28.3.17 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
```

166 Class Documentation

7.28.3.18 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
( )
```

7.28.3.19 reset_array() [2/2]

7.28.3.20 set_counters()

7.28.3.21 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_ )
```

7.28.3.22 set_rules_dyn()

7.28.4 Member Data Documentation

7.28.4.1 change_stats

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector< double > > Support< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::change_stats
```

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

7.28.4.2 coordinates free

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::pair<uint,uint> > Support< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::coordinates_free
```

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

7.28.4.3 coordinates locked

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::pair<uint,uint> > Support< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::coordinates_locked
```

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

7.28.4.4 current_stats

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

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

7.28.4.5 delete_counters

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

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

168 Class Documentation

7.28.4.6 delete_rules

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

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

7.28.4.7 delete_rules_dyn

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

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

7.28.4.8 M

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
uint Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::M
```

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

7.28.4.9 max_num_elements

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
uint Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::max_num_←
elements = BARRY_MAX_NUM_ELEMENTS
```

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

7.28.4.10 N

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
uint Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::N
```

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

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

• include/barry/support-bones.hpp

7.29 vecHasher< T > Struct Template Reference

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

Public Member Functions

• std::size_t operator() (std::vector< T > const &dat) const noexcept

7.29.1 Detailed Description

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

Definition at line 86 of file typedefs.hpp.

7.29.2 Member Function Documentation

7.29.2.1 operator()()

Definition at line 87 of file typedefs.hpp.

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

• include/barry/typedefs.hpp

170 Class Documentation

Chapter 8

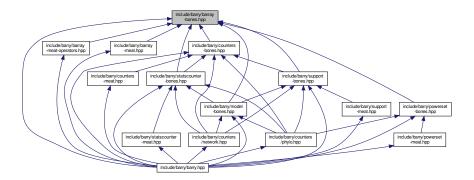
File Documentation

8.1 include/barry/barray-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "cell-bones.hpp"
#include "barraycell-bones.hpp"
Include dependency graph for barray-bones.hpp:
```



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



Classes

class BArray < Cell_Type, Data_Type >
 Baseline class for binary arrays.

Macros

• #define BARRAY_BONES_HPP 1

8.1.1 Macro Definition Documentation

8.1.1.1 BARRAY_BONES_HPP

#define BARRAY_BONES_HPP 1

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

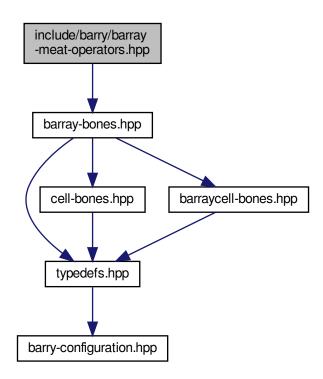
8.2 include/barry/barray-iterator.hpp File Reference

Classes

class ConstBArrayRowIter< Cell_Type, Data_Type >

8.3 include/barry/barray-meat-operators.hpp File Reference

#include "barray-bones.hpp"
Include dependency graph for barray-meat-operators.hpp:



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



Macros

- #define BARRY_BARRAY_MEAT_OPERATORS_HPP 1
- #define ROW(a) this->el_ij[a]
- #define COL(a) this->el_ji[a]

Functions

template<typename Cell_Type , typename Data_Type >
 void checkdim_ (const BArray< Cell_Type, Data_Type > &lhs, const BArray< Cell_Type, Data_Type > &rhs)

8.3.1 Macro Definition Documentation

8.3.1.1 BARRY_BARRAY_MEAT_OPERATORS_HPP

```
#define BARRY_BARRAY_MEAT_OPERATORS_HPP 1
```

Definition at line 5 of file barray-meat-operators.hpp.

8.3.1.2 COL

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

8.3.1.3 ROW

Definition at line 7 of file barray-meat-operators.hpp.

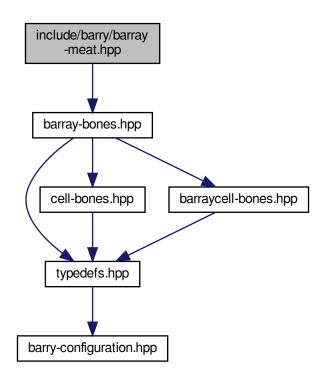
8.3.2 Function Documentation

8.3.2.1 checkdim_()

Definition at line 11 of file barray-meat-operators.hpp.

8.4 include/barry/barray-meat.hpp File Reference

#include "barray-bones.hpp"
Include dependency graph for barray-meat.hpp:



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



Macros

- #define ROW(a) this->el_ij[a]
- #define COL(a) this->el_ji[a]

8.4.1 Macro Definition Documentation

8.4.1.1 COL

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

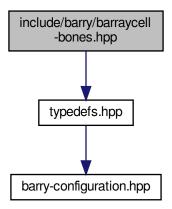
8.4.1.2 ROW

```
#define ROW( a \ ) \ {\tt this->el_ij[a]}
```

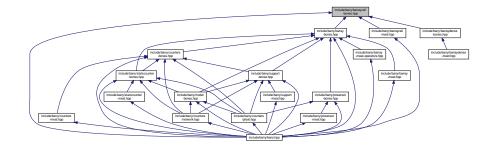
Definition at line 7 of file barray-meat.hpp.

8.5 include/barry/barraycell-bones.hpp File Reference

```
#include "typedefs.hpp"
Include dependency graph for barraycell-bones.hpp:
```



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

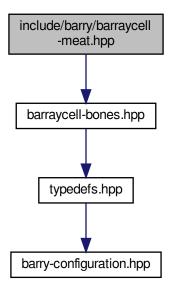


Classes

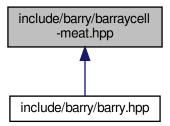
- class BArrayCell
 Cell_Type, Data_Type
- class BArrayCell_const< Cell_Type, Data_Type >

8.6 include/barry/barraycell-meat.hpp File Reference

#include "barraycell-bones.hpp"
Include dependency graph for barraycell-meat.hpp:



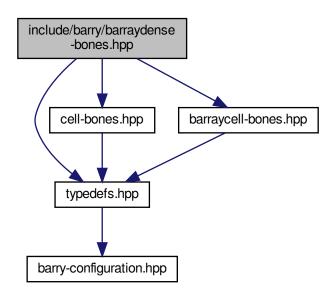
This graph shows which files directly or indirectly include this file:



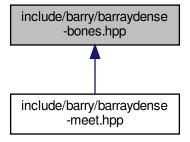
8.7 include/barry/barraydense-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "cell-bones.hpp"
#include "barraycell-bones.hpp"
```

Include dependency graph for barraydense-bones.hpp:



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



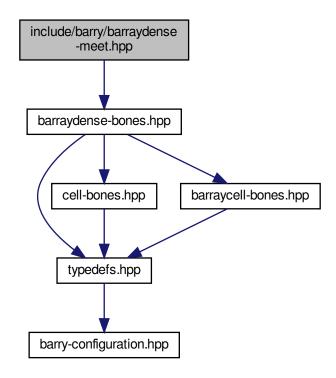
Classes

class BArrayDense < Cell_Type, Data_Type >
 Baseline class for binary arrays.

8.8 include/barry/barraydense-meet.hpp File Reference

#include "barraydense-bones.hpp"

Include dependency graph for barraydense-meet.hpp:



Macros

- #define BARRY_BARRAYDENSE_MEAT_HPP
- #define ROW(a) this->el_ij[a]
- #define COL(a) this->el_ji[a]
- #define POS(a, b) (b)*N + (a)
- #define ZERO_CELL Cell< Cell_Type >(static_cast< Cell_Type >(0.0))

8.8.1 Macro Definition Documentation

8.8.1.1 BARRY_BARRAYDENSE_MEAT_HPP

#define BARRY_BARRAYDENSE_MEAT_HPP

Definition at line 5 of file barraydense-meet.hpp.

8.8.1.2 COL

```
#define COL( a \ ) \ \  \  this->el_ji[a]
```

Definition at line 8 of file barraydense-meet.hpp.

8.8.1.3 POS

```
#define POS(  a, \\ b ) \ (b)*N + (a)
```

Definition at line 9 of file barraydense-meet.hpp.

8.8.1.4 ROW

```
#define ROW( a \ ) \ {\tt this->el_ij[a]}
```

Definition at line 7 of file barraydense-meet.hpp.

8.8.1.5 ZERO_CELL

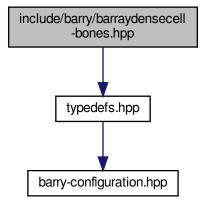
```
#define ZERO_CELL Cell Cell_Type >(static_cast< Cell_Type >(0.0))
```

Definition at line 14 of file barraydense-meet.hpp.

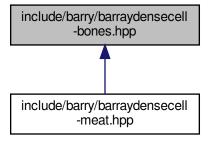
8.9 include/barry/barraydensecell-bones.hpp File Reference

```
#include "typedefs.hpp"
```

Include dependency graph for barraydensecell-bones.hpp:



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

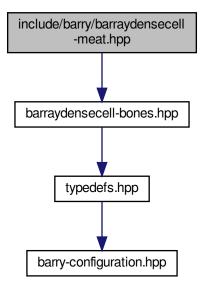


Classes

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

8.10 include/barry/barraydensecell-meat.hpp File Reference

#include "barraydensecell-bones.hpp"
Include dependency graph for barraydensecell-meat.hpp:



Macros

- #define BARRY_BARRAYDENSECELL_MEAT_HPP 1
- #define POS(a, b) (a) + (b) * Array->N

8.10.1 Macro Definition Documentation

8.10.1.1 BARRY_BARRAYDENSECELL_MEAT_HPP

```
#define BARRY_BARRAYDENSECELL_MEAT_HPP 1
```

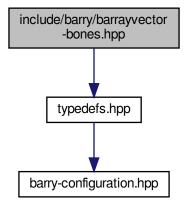
Definition at line 4 of file barraydensecell-meat.hpp.

8.10.1.2 POS

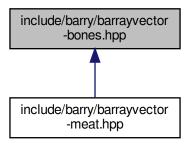
Definition at line 6 of file barraydensecell-meat.hpp.

8.11 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:

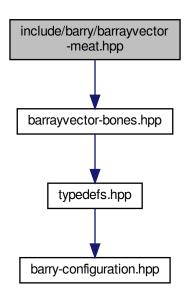


Classes

- class BArrayVector < Cell_Type, Data_Type >
 Row or column of a BArray
- class BArrayVector_const < Cell_Type, Data_Type >

8.12 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

8.12.1 Macro Definition Documentation

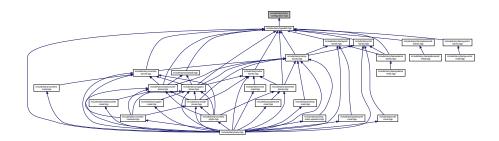
8.12.1.1 BARRY BARRAYVECTOR MEAT HPP

#define BARRY_BARRAYVECTOR_MEAT_HPP 1

Definition at line 4 of file barrayvector-meat.hpp.

8.13 include/barry/barry-configuration.hpp File Reference

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



Configuration MACROS

These are mostly related to performance. The definitions follow:

- BARRY_USE_UNORDERED_MAP If specified, then barry is compiled using std::unordered_map. Otherwise it will use std::map for the arrays.
- BARRY_USE_SAFE_EXP When specified, it will multiply all likelihoods in Model by (1/-100)/(1/-100) so that numerical overflows are avoided.
- BARRY_USE_ISFINITE When specified, it will introduce a macro that checks whether the likelihood is finite or not.
- printf_barry If not specified, will be defined as printf.
- #define BARRY_SAFE_EXP -100.0
- #define BARRY_ISFINITE(a)
- #define BARRY CHECK SUPPORT(x, maxs)
- #define printf barry printf
- #define BARRY_MAX_NUM_ELEMENTS static_cast< unsigned int >(UINT_MAX/2u)
- template < typename Ta , typename Tb > using Map = std::map < Ta, Tb >

8.13.1 Macro Definition Documentation

8.13.1.1 BARRY_CHECK_SUPPORT

```
#define BARRY_CHECK_SUPPORT(
          x,
          maxs )
```

Definition at line 45 of file barry-configuration.hpp.

8.13.1.2 BARRY_ISFINITE

Definition at line 38 of file barry-configuration.hpp.

8.13.1.3 BARRY_MAX_NUM_ELEMENTS

```
#define BARRY_MAX_NUM_ELEMENTS static_cast< unsigned int >(UINT_MAX/2u)
```

Definition at line 53 of file barry-configuration.hpp.

8.13.1.4 BARRY_SAFE_EXP

```
#define BARRY_SAFE_EXP -100.0
```

Definition at line 31 of file barry-configuration.hpp.

8.13.1.5 printf_barry

```
#define printf_barry printf
```

Definition at line 49 of file barry-configuration.hpp.

8.13.2 Typedef Documentation

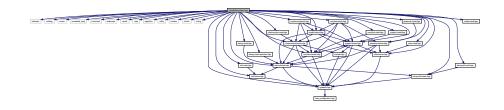
8.13.2.1 Map

```
template<typename Ta , typename Tb >
using Map = std::map<Ta,Tb>
```

Definition at line 25 of file barry-configuration.hpp.

8.14 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 <string>
#include "typedefs.hpp"
#include "cell-bones.hpp"
#include "cell-meat.hpp"
#include "barray-bones.hpp"
#include "barraycell-bones.hpp"
#include "barray-meat.hpp"
#include "barraycell-meat.hpp"
#include "barray-meat-operators.hpp"
#include "counters-bones.hpp"
#include "counters-meat.hpp"
#include "statscounter-bones.hpp"
#include "statscounter-meat.hpp"
#include "support-bones.hpp"
#include "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 dependency graph for barry.hpp:
```



Namespaces

barry

barry: Your go-to motif accountant

· barry::counters

Tree class and Treelterator class.

- barry::counters::network
- · barry::counters::phylo

Macros

- #define BARRY_HPP
- #define BARRY_VERSION 0.1
- #define COUNTER FUNCTION(a)
- #define COUNTER_LAMBDA(a)
- #define RULE_FUNCTION(a)
- #define RULE_LAMBDA(a)

8.14.1 Macro Definition Documentation

8.14.1.1 BARRY_HPP

```
#define BARRY_HPP
```

Definition at line 20 of file barry.hpp.

8.14.1.2 BARRY_VERSION

```
#define BARRY_VERSION 0.1
```

Definition at line 22 of file barry.hpp.

8.14.1.3 COUNTER_FUNCTION

```
#define COUNTER_FUNCTION( a )
```

Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
inline double (a) (const Array_Type & Array, uint i, uint j, Data_Type * data) \
```

Definition at line 73 of file barry.hpp.

8.14.1.4 COUNTER_LAMBDA

Definition at line 76 of file barry.hpp.

8.14.1.5 RULE_FUNCTION

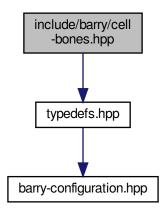
Definition at line 80 of file barry.hpp.

Definition at line 83 of file barry.hpp.

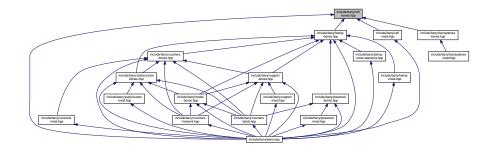
8.14.1.6 RULE_LAMBDA

8.15 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:

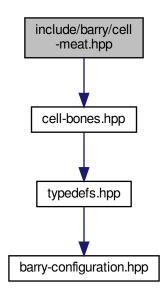


Classes

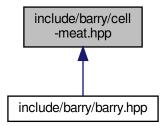
class Cell
 Cell_Type >
 Entries in BArray. For now, it only has two members:

8.16 include/barry/cell-meat.hpp File Reference

#include "cell-bones.hpp"
Include dependency graph for cell-meat.hpp:



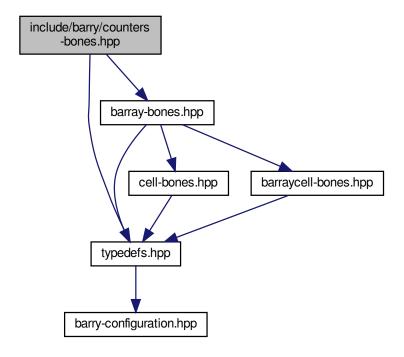
This graph shows which files directly or indirectly include this file:



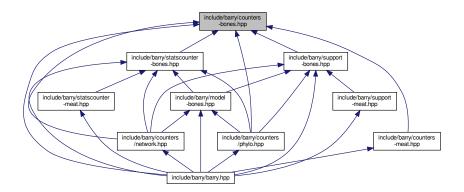
8.17 include/barry/col-bones.hpp File Reference

8.18 include/barry/counters-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "barray-bones.hpp"
Include dependency graph for counters-bones.hpp:
```



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



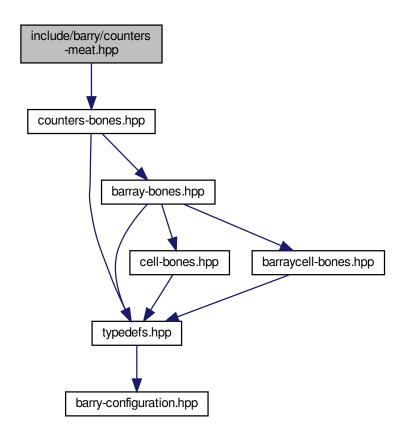
Classes

- class Counter < Array_Type, Data_Type >
 A counter function based on change statistics.

8.19 include/barry/counters-meat.hpp File Reference

#include "counters-bones.hpp"

Include dependency graph for counters-meat.hpp:



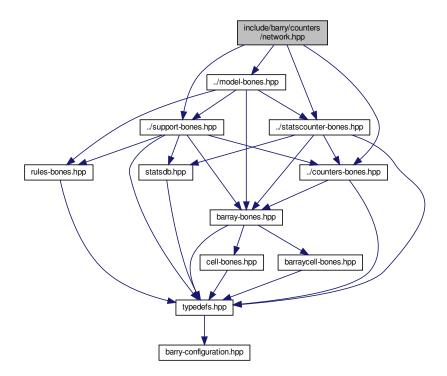
This graph shows which files directly or indirectly include this file:



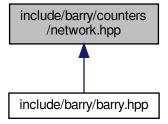
8.20 include/barry/counters/network.hpp File Reference

```
#include "../counters-bones.hpp"
#include "../support-bones.hpp"
```

```
#include "../statscounter-bones.hpp"
#include "../model-bones.hpp"
Include dependency graph for network.hpp:
```



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



Classes

class NetworkData

Data class for Networks.

· class NetCounterData

Data class used to store arbitrary uint or double vectors.

Macros

- #define NET_C_DATA_IDX(i) (data->indices[i])
- #define NET_C_DATA_NUM(i) (data->numbers[i])

Macros for defining counters

- #define NETWORK COUNTER(a)
- #define NETWORK_COUNTER_LAMBDA(a)

Macros for defining rules

- #define NETWORK_RULE(a)
- #define NETWORK_RULE_LAMBDA(a)

Typedefs

Convenient typedefs for network objects.

- typedef BArray< double, NetworkData > Network
- typedef Counter
 NetWork, NetCounterData > NetCounter
- typedef Counters < Network, NetCounterData > NetCounters
- typedef Support < Network, NetCounterData > NetSupport
- typedef StatsCounter< Network, NetCounterData > NetStatsCounter
- typedef Model < Network, NetCounterData > NetModel
- typedef Rule < Network, bool > NetRule
- typedef Rules < Network, bool > NetRules

Functions

• void counter_edges (NetCounters *counters)

Number of edges.

void counter_isolates (NetCounters *counters)

Number of isolated vertices.

void counter_mutual (NetCounters *counters)

Number of mutual ties.

- void counter_istar2 (NetCounters *counters)
- void counter ostar2 (NetCounters *counters)
- void counter_ttriads (NetCounters *counters)
- void counter ctriads (NetCounters *counters)
- void counter_density (NetCounters *counters)
- void counter_idegree15 (NetCounters *counters)
- void counter odegree15 (NetCounters *counters)
- void counter_absdiff (NetCounters *counters, uint attr_id, double alpha=1.0)

Sum of absolute attribute difference between ego and alter.

• void counter_diff (NetCounters *counters, uint attr_id, double alpha=1.0, double tail_head=true)

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

- NETWORK_COUNTER (init_single_attr)
- void counter_nodeicov (NetCounters *counters, uint attr_id)
- void counter_nodeocov (NetCounters *counters, uint attr_id)

- void counter_nodecov (NetCounters *counters, uint attr_id)
- void counter nodematch (NetCounters *counters, uint attr id)
- void counter_idegree (NetCounters *counters, std::vector< uint > d)

Counts number of vertices with a given in-degree.

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

Counts number of vertices with a given out-degree.

void counter_degree (NetCounters *counters, std::vector< uint > d)

Counts number of vertices with a given out-degree.

void counter_css_partially_false_recip (NetCounters *counters, uint netsize, const std::vector< uint > &end)

Counts errors of commission.

Rules for network models

Parameters

```
rules | A pointer to a NetRules object (Rules < Network, bool > ).
```

void rules_zerodiag (NetRules *rules)
 Number of edges.

8.20.1 Macro Definition Documentation

8.20.1.1 NET_C_DATA_IDX

Definition at line 79 of file network.hpp.

8.20.1.2 NET_C_DATA_NUM

Definition at line 80 of file network.hpp.

8.20.1.3 NETWORK_COUNTER

Function for definition of a network counter function

Definition at line 101 of file network.hpp.

8.20.1.4 NETWORK_COUNTER_LAMBDA

Lambda function for definition of a network counter function

Definition at line 104 of file network.hpp.

8.20.1.5 NETWORK_RULE

Function for definition of a network counter function

Definition at line 113 of file network.hpp.

8.20.1.6 NETWORK_RULE_LAMBDA

Lambda function for definition of a network counter function

Definition at line 116 of file network.hpp.

8.20.2 Typedef Documentation

8.20.2.1 NetCounter

typedef Counter<Network, NetCounterData > NetCounter

Definition at line 88 of file network.hpp.

8.20.2.2 NetCounters

typedef Counters< Network, NetCounterData> NetCounters

Definition at line 89 of file network.hpp.

8.20.2.3 NetModel

typedef Model < Network, NetCounterData > NetModel

Definition at line 92 of file network.hpp.

8.20.2.4 NetRule

typedef Rule<Network,bool> NetRule

Definition at line 93 of file network.hpp.

8.20.2.5 NetRules

typedef Rules<Network,bool> NetRules

Definition at line 94 of file network.hpp.

8.20.2.6 NetStatsCounter

```
typedef StatsCounter<Network, NetCounterData> NetStatsCounter
```

Definition at line 91 of file network.hpp.

8.20.2.7 NetSupport

```
typedef Support<Network, NetCounterData > NetSupport
```

Definition at line 90 of file network.hpp.

8.20.2.8 Network

```
typedef BArray<double, NetworkData> Network
```

Definition at line 87 of file network.hpp.

8.20.3 Function Documentation

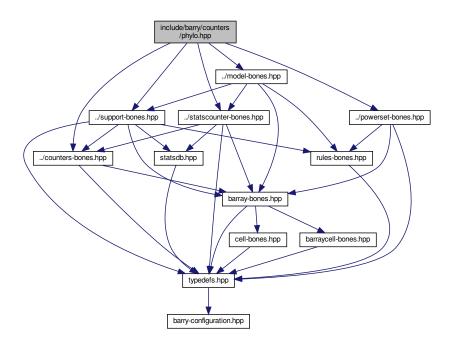
8.20.3.1 rules_zerodiag()

Number of edges.

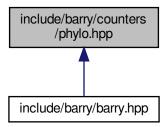
Definition at line 909 of file network.hpp.

8.21 include/barry/counters/phylo.hpp File Reference

```
#include "../counters-bones.hpp"
#include "../support-bones.hpp"
#include "../statscounter-bones.hpp"
#include "../model-bones.hpp"
#include "../powerset-bones.hpp"
Include dependency graph for phylo.hpp:
```



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



Classes

class NodeData

Data definition for the PhyloArray class.

- class PhyloCounterData
- class PhyloRuleDynData

Macros

- #define PHYLO COUNTER LAMBDA(a)
 - Extension of a simple counter.
- #define PHYLO RULE DYN LAMBDA(a)
- #define PHYLO_CHECK_MISSING()

Typedefs

typedef std::vector< std::pair< uint, uint > > PhyloRuleData

Convenient typedefs for Node objects.

- typedef BArray< uint, NodeData > PhyloArray
- typedef Counter< PhyloArray, PhyloCounterData > PhyloCounter
- typedef Counters
 PhyloArray, PhyloCounterData > PhyloCounters
- typedef Rule < PhyloArray, PhyloRuleData > PhyloRule
- typedef Rules
 PhyloArray, PhyloRuleData
 PhyloRules
- typedef Rule
 PhyloArray, PhyloRuleDynData
 PhyloRuleDyn
- typedef Rules
 PhyloArray, PhyloRuleDynData
 PhyloRulesDyn
- typedef Support < PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloSupport
- typedef StatsCounter< PhyloArray, PhyloCounterData > PhyloStatsCounter
- typedef Model < PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloModel
- typedef PowerSet< PhyloArray, PhyloRuleData > PhyloPowerSet

Functions

- std::string get last name (bool d)
- void counter_overall_gains (PhyloCounters *counters, bool duplication=true)

Overall functional gains.

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

k genes gain function nfun

• void counter_genes_changing (PhyloCounters *counters, bool duplication=true)

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

void counter_prop_genes_changing (PhyloCounters *counters, bool duplication=true)

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

void counter overall loss (PhyloCounters *counters, bool duplication=true)

Overall functional loss.

• void counter_maxfuns (PhyloCounters *counters, uint lb, uint ub, bool duplication=true)

Cap the number of functions per gene.

void counter loss (PhyloCounters *counters, std::vector< uint > nfun, bool duplication=true)

Total count of losses for an specific function.

• void counter_overall_changes (PhyloCounters *counters, bool duplication=true)

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

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

Total count of Sub-functionalization events.

• void counter_cogain (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)

Co-evolution (joint gain or loss)

void counter_longest (PhyloCounters *counters)

Longest branch mutates (either by gain or by loss)

- void counter_neofun (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)
 - Total number of neofunctionalization events.
- void counter_neofun_a2b (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)

 Total number of neofunctionalization events.
- void counter_co_opt (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)
 Function co-opting.
- void rule_dyn_limit_changes (PhyloSupport *support, uint pos, uint lb, uint ub, bool duplication=true)

 Overall functional gains.

8.21.1 Macro Definition Documentation

8.21.1.1 PHYLO CHECK MISSING

```
#define PHYLO_CHECK_MISSING( )
```

Value:

```
if (Array.D() == nullptr) \
throw std::logic_error("The array data is nullptr."); \
if (data == nullptr) \
throw std::logic_error("The counter/rule data is nullptr.")
```

Definition at line 121 of file phylo.hpp.

8.21.1.2 PHYLO_COUNTER_LAMBDA

Value:

```
Counter_fun_type<PhyloArray, PhyloCounterData> a = \
[](const PhyloArray & Array, uint i, uint j, PhyloCounterData * data)
```

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 115 of file phylo.hpp.

8.21.1.3 PHYLO_RULE_DYN_LAMBDA

Value:

```
Rule_fun_type<PhyloArray, PhyloRuleDynData> a = \
[](const PhyloArray & Array, uint i, uint j, PhyloRuleDynData * data)
```

Definition at line 118 of file phylo.hpp.

8.21.2 Typedef Documentation

8.21.2.1 PhyloArray

```
typedef BArray<uint, NodeData> PhyloArray
```

Definition at line 88 of file phylo.hpp.

8.21.2.2 PhyloCounter

```
typedef Counter<PhyloArray, PhyloCounterData > PhyloCounter
```

Definition at line 89 of file phylo.hpp.

8.21.2.3 PhyloCounters

```
typedef Counters< PhyloArray, PhyloCounterData> PhyloCounters
```

Definition at line 90 of file phylo.hpp.

8.21.2.4 PhyloModel

```
{\tt typedef\ Model < PhyloArray,\ PhyloCounterData,\ PhyloRuleData,\ PhyloRuleDynData > PhyloModel}
```

Definition at line 100 of file phylo.hpp.

8.21.2.5 PhyloPowerSet

typedef PowerSet<PhyloArray, PhyloRuleData> PhyloPowerSet

Definition at line 101 of file phylo.hpp.

8.21.2.6 PhyloRule

typedef Rule<PhyloArray,PhyloRuleData> PhyloRule

Definition at line 92 of file phylo.hpp.

8.21.2.7 PhyloRuleData

typedef std::vector< std::pair< uint, uint > > PhyloRuleData

Definition at line 81 of file phylo.hpp.

8.21.2.8 PhyloRuleDyn

typedef Rule<PhyloArray,PhyloRuleDynData> PhyloRuleDyn

Definition at line 95 of file phylo.hpp.

8.21.2.9 PhyloRules

typedef Rules<PhyloArray,PhyloRuleData> PhyloRules

Definition at line 93 of file phylo.hpp.

8.21.2.10 PhyloRulesDyn

typedef Rules<PhyloArray,PhyloRuleDynData> PhyloRulesDyn

Definition at line 96 of file phylo.hpp.

8.21.2.11 PhyloStatsCounter

typedef StatsCounter<PhyloArray, PhyloCounterData> PhyloStatsCounter Definition at line 99 of file phylo.hpp.

8.21.2.12 PhyloSupport

typedef Support<PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloSupport Definition at line 98 of file phylo.hpp.

8.21.3 Function Documentation

8.21.3.1 get_last_name()

```
std::string get_last_name (
          bool d ) [inline]
```

Definition at line 126 of file phylo.hpp.

8.22 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:
```

support-bones.hpp statscounter-bones.hpp

rules-bones.hpp counters-bones.hpp

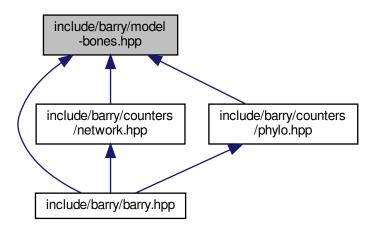
barray-bones.hpp

barraycell-bones.hpp

typedefs.hpp

barry-configuration.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)

8.22.1 Function Documentation

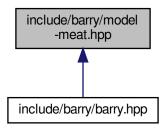
8.22.1.1 keygen_default()

Array Hasher class (used for computing support)

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

8.23 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 std::vector< double > ¶ms, const Counts_type &support)
- double likelihood_ (const std::vector< double > &target_stats, const std::vector< double > ¶ms, const double normalizing_constant, bool log_=false)
- MODEL_TEMPLATE (, Model)()
- MODEL_TEMPLATE (, Model)(const MODEL_TYPE() &Model_)

8.23.1 Macro Definition Documentation

8.23.1.1 MODEL_TEMPLATE

Definition at line 75 of file model-meat.hpp.

8.23.1.2 MODEL_TEMPLATE_ARGS

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

8.23.1.3 MODEL_TYPE

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

8.23.2 Function Documentation

8.23.2.1 likelihood_()

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

8.23.2.2 MODEL_TEMPLATE() [1/2]

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

8.23.2.3 MODEL_TEMPLATE() [2/2]

Definition at line 134 of file model-meat.hpp.

8.23.2.4 update_normalizing_constant()

Definition at line 11 of file model-meat.hpp.

8.24 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-meet.hpp"
Include dependency graph for geese.hpp:
```



8.25 include/barry/models/geese/flock-bones.hpp File Reference

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

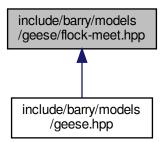


Classes

· class Flock

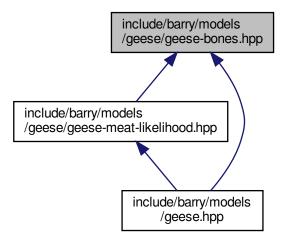
A Flock is a group of Geese.

8.26 include/barry/models/geese/flock-meet.hpp File Reference



8.27 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)
- $\bullet \ \ \mathsf{bool} \ \mathsf{vec_diff} \ (\mathsf{const} \ \mathsf{std} : \mathsf{vector} < \mathsf{unsigned} \ \mathsf{int} > \mathsf{\&s}, \ \mathsf{const} \ \mathsf{std} : \mathsf{vector} < \mathsf{unsigned} \ \mathsf{int} > \mathsf{\&a}) \\$

8.27.1 Macro Definition Documentation

8.27.1.1 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.

8.27.2 Function Documentation

8.27.2.1 keygen_full()

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

8.27.2.2 RULE_FUNCTION()

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

8.27.2.3 vec_diff()

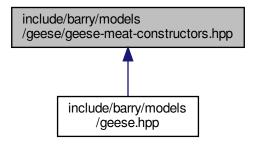
Definition at line 59 of file geese-bones.hpp.

8.27.2.4 vector_caster()

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

8.28 include/barry/models/geese/geese-meat-constructors.hpp File Reference

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



8.29 include/barry/models/geese/geese-meat-likelihood.hpp File Reference

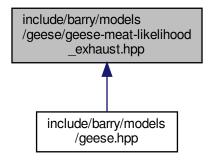
#include "geese-bones.hpp"
Include dependency graph for geese-meat-likelihood.hpp:



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



8.30 include/barry/models/geese/geese-meat-likelihood_exhaust.hpp File Reference

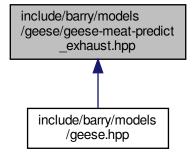


8.31 include/barry/models/geese/geese-meat-predict.hpp File Reference

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



8.32 include/barry/models/geese/geese-meat-predict_exhaust.hpp File Reference



8.33 include/barry/models/geese/geese-meat-predict_sim.hpp File Reference

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



8.34 include/barry/models/geese/geese-meat-simulate.hpp File Reference



8.35 include/barry/models/geese/geese-meat.hpp File Reference

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



8.36 include/barry/models/geese/geese-node-bones.hpp File Reference

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



Classes

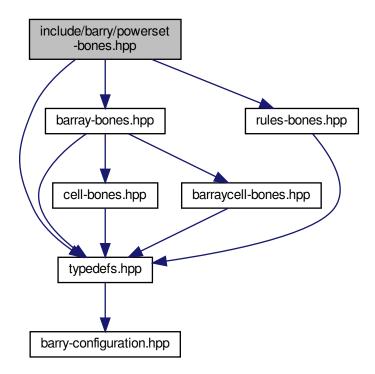
• class Node

A single node for the model.

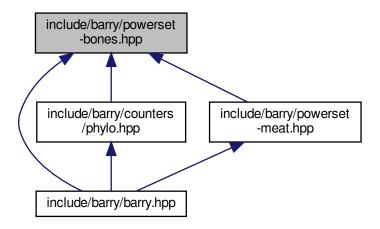
8.37 include/barry/powerset-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "barray-bones.hpp"
#include "rules-bones.hpp"
```

Include dependency graph for powerset-bones.hpp:



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

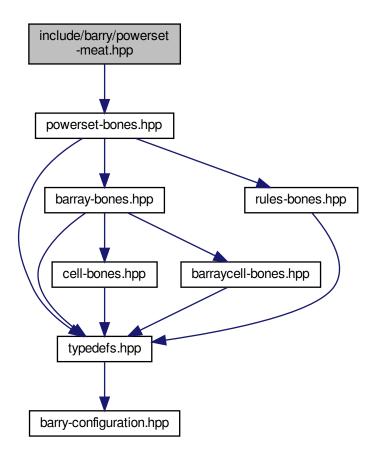


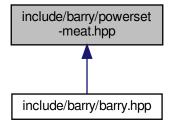
Classes

8.38 include/barry/powerset-meat.hpp File Reference

#include "powerset-bones.hpp"

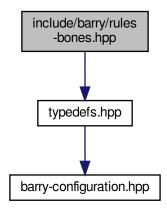
Include dependency graph for powerset-meat.hpp:



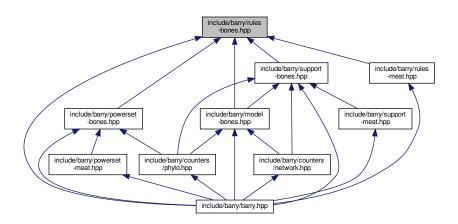


8.39 include/barry/rules-bones.hpp File Reference

#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.
- $\bullet \ \ {\it class Rules} {< Array_Type, Data_Type} >$

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)

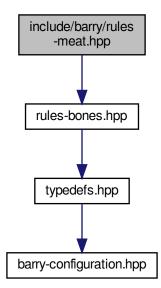
8.39.1 Function Documentation

8.39.1.1 rule_fun_default()

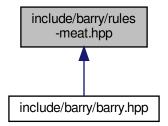
Definition at line 10 of file rules-bones.hpp.

8.40 include/barry/rules-meat.hpp File Reference

```
#include "rules-bones.hpp"
Include dependency graph for rules-meat.hpp:
```



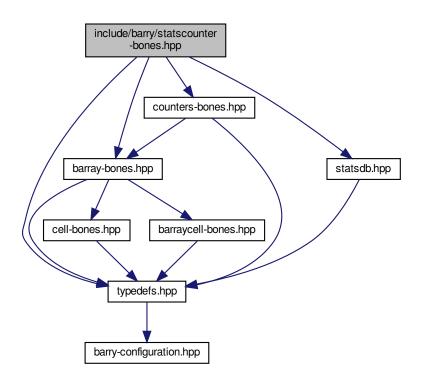
This graph shows which files directly or indirectly include this file:



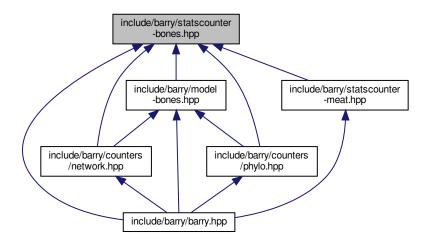
8.41 include/barry/statscounter-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "barray-bones.hpp"
#include "statsdb.hpp"
#include "counters-bones.hpp"
```

Include dependency graph for statscounter-bones.hpp:



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



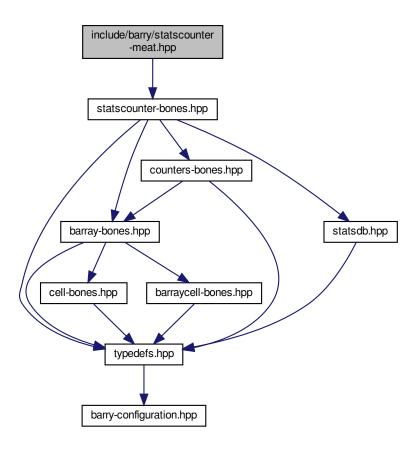
Classes

class StatsCounter< Array_Type, Data_Type >
 Count stats for a single Array.

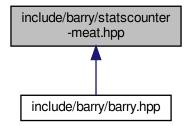
8.42 include/barry/statscounter-meat.hpp File Reference

#include "statscounter-bones.hpp"

Include dependency graph for statscounter-meat.hpp:



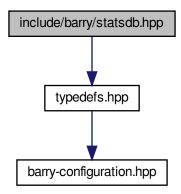
This graph shows which files directly or indirectly include this file:



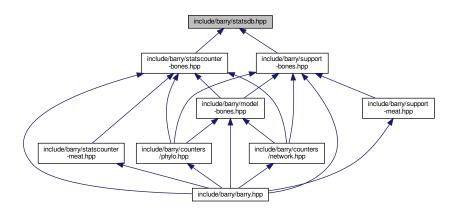
8.43 include/barry/statsdb.hpp File Reference

#include "typedefs.hpp"

Include dependency graph for statsdb.hpp:



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



Classes

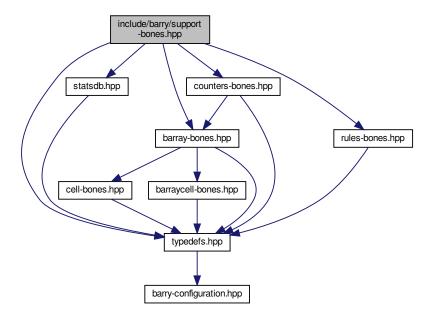
class FreqTable < T >

Database of statistics.

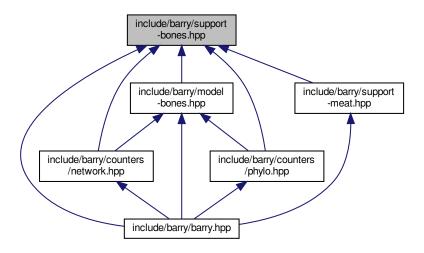
8.44 include/barry/support-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "barray-bones.hpp"
#include "statsdb.hpp"
#include "counters-bones.hpp"
```

#include "rules-bones.hpp"
Include dependency graph for support-bones.hpp:



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

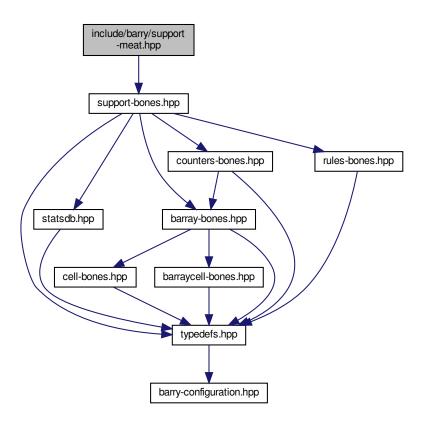


Classes

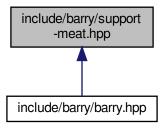
class Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >
 Compute the support of sufficient statistics.

8.45 include/barry/support-meat.hpp File Reference

#include "support-bones.hpp"
Include dependency graph for support-meat.hpp:



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



Macros

• #define BARRY_SUPPORT_MEAT_HPP 1

- #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)(uint pos
- calc_backend (pos+1u, array_bank, stats_bank)
- EmptyArray insert_cell (cfree.first, cfree.second, EmptyArray.default_val().value, false, false)
- for (uint n=0u;n< counters->size();++n) = change_stats[pos][n]
- if (rules_dyn->size() > 0u)
- if (array bank !=nullptr) array bank -> push back(EmptyArray)
- if (stats_bank !=nullptr) stats_bank -> push_back(current_stats)
- EmptyArray rm_cell (cfree.first, cfree.second, 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 (Counts_type, get_counts)() const
- SUPPORT_TEMPLATE (const MapVec_type<> *, get_counts_ptr)() 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< std::vector< double > > * stats_bank
- const std::pair < uint, uint > & cfree = coordinates free[pos]
- else
- 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_

8.45.1 Macro Definition Documentation

8.45.1.1 BARRY_SUPPORT_MEAT_HPP

```
#define BARRY_SUPPORT_MEAT_HPP 1
```

Definition at line 4 of file support-meat.hpp.

8.45.1.2 SUPPORT_TEMPLATE

Value:

```
template SUPPORT_TEMPLATE_ARGS() \
inline a SUPPORT_TYPE()::b
```

Definition at line 12 of file support-meat.hpp.

8.45.1.3 SUPPORT_TEMPLATE_ARGS

```
template SUPPORT_TEMPLATE_ARGS( )
```

Value:

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

8.45.1.4 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.

8.45.2 Function Documentation

8.45.2.1 calc_backend()

```
calc_backend (
    pos+ 1u,
    array_bank ,
    stats_bank )
```

8.45.2.2 for()

```
for (  \label{eq:uint_n} \mbox{uint } n = 0 \mbox{$u$;} \mbox{$n$< $counters-$>$size()$; ++$n ) = $change\_stats[pos][n]
```

Definition at line 134 of file support-meat.hpp.

8.45.2.3 if() [1/3]

```
if (
     array_bank ! = nullptr ) -> push_back(EmptyArray)
```

8.45.2.4 if() [2/3]

```
if (
    rules_dyn-> size(),
    0u )
```

Definition at line 148 of file support-meat.hpp.

8.45.2.5 if() [3/3]

```
if (
     stats_bank ! = nullptr ) -> push_back(current_stats)
```

8.45.2.6 insert_cell()

8.45.2.7 rm_cell()

8.45.2.8 **SUPPORT_TEMPLATE()** [1/17]

```
SUPPORT_TEMPLATE (
          bool ,
          eval_rules_dyn ) const
```

Definition at line 320 of file support-meat.hpp.

8.45.2.9 SUPPORT_TEMPLATE() [2/17]

Definition at line 369 of file support-meat.hpp.

8.45.2.10 SUPPORT_TEMPLATE() [3/17]

Definition at line 348 of file support-meat.hpp.

8.45.2.11 SUPPORT_TEMPLATE() [4/17]

```
SUPPORT_TEMPLATE (

Counts_type ,

get_counts ) const
```

Definition at line 342 of file support-meat.hpp.

8.45.2.12 SUPPORT_TEMPLATE() [5/17]

```
SUPPORT_TEMPLATE (
          std::vector< double > * ,
          get_current_stats )
```

Definition at line 354 of file support-meat.hpp.

8.45.2.13 SUPPORT_TEMPLATE() [6/17]

```
SUPPORT_TEMPLATE (
     void ,
     add_counter )
```

8.45.2.14 SUPPORT_TEMPLATE() [7/17]

```
SUPPORT_TEMPLATE (
     void ,
     add_rule )
```

8.45.2.15 SUPPORT_TEMPLATE() [8/17]

8.45.2.16 SUPPORT_TEMPLATE() [9/17]

```
SUPPORT_TEMPLATE ( void , calc )
```

Definition at line 197 of file support-meat.hpp.

8.45.2.17 SUPPORT_TEMPLATE() [10/17]

8.45.2.18 SUPPORT_TEMPLATE() [11/17]

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

8.45.2.19 SUPPORT_TEMPLATE() [12/17]

```
SUPPORT_TEMPLATE (
     void ,
     print ) const
```

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

8.45.2.20 SUPPORT_TEMPLATE() [13/17]

```
SUPPORT_TEMPLATE (
     void ,
     reset_array )
```

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

8.45.2.21 SUPPORT_TEMPLATE() [14/17]

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

8.45.2.22 SUPPORT_TEMPLATE() [15/17]

8.45.2.23 SUPPORT_TEMPLATE() [16/17]

```
SUPPORT_TEMPLATE (
     void ,
     set_rules )
```

8.45.2.24 SUPPORT_TEMPLATE() [17/17]

8.45.3 Variable Documentation

8.45.3.1 array_bank

```
std::vector< Array_Type >* array_bank
```

Definition at line 109 of file support-meat.hpp.

8.45.3.2 cfree

```
const std::pair<uint,uint>& cfree = coordinates_free[pos]
```

Definition at line 123 of file support-meat.hpp.

8.45.3.3 counters

```
counters = counters_
```

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

8.45.3.4 counters_

```
Data_Counter_Type* counters_
Initial value:
{
    if (delete_counters)
        delete counters
```

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

8.45.3.5 delete_counters

```
delete_counters = false
```

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

8.45.3.6 delete_rules

```
delete_rules = false
```

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

8.45.3.7 delete_rules_dyn

```
delete\_rules\_dyn = false
```

Definition at line 313 of file support-meat.hpp.

8.45.3.8 else

else

Initial value:

```
data.add(current_stats)
```

Definition at line 166 of file support-meat.hpp.

8.45.3.9 f_

```
Data_Rule_Dyn_Type f_
Initial value:
{
    counters->add_counter(f_)
```

Definition at line 223 of file support-meat.hpp.

8.45.3.10 return

return

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

8.45.3.11 rules

```
rules = rules_
```

Definition at line 282 of file support-meat.hpp.

8.45.3.12 rules_

```
Data_Rule_Dyn_Type * rules_
Initial value:
{
    if (delete_rules)
        delete rules
```

Definition at line 275 of file support-meat.hpp.

8.45.3.13 rules_dyn

```
rules_dyn = rules_
```

Definition at line 314 of file support-meat.hpp.

8.45.3.14 stats_bank

```
std::vector< Array_Type > std::vector< std::vector< double > >* stats_bank
```

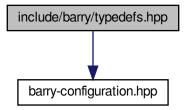
Initial value:

```
if (pos >= coordinates_free.size())
    return
```

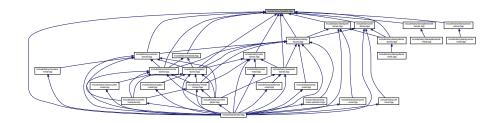
Definition at line 110 of file support-meat.hpp.

8.46 include/barry/typedefs.hpp File Reference

#include "barry-configuration.hpp"
Include dependency graph for typedefs.hpp:



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



Classes

- class Entries < Cell_Type >
 - A wrapper class to store source, target, val from a BArray object.
- struct vecHasher< T >

Namespaces

CHECK

Integer constants used to specify which cell should be check.

• EXISTS

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

Typedefs

```
• typedef unsigned int uint
```

```
    typedef std::vector< std::pair< std::vector< double >, uint > > Counts_type
```

```
    template<typename Cell_Type >
        using Row_type = Map< uint, Cell< Cell_Type > >
    template<typename Cell_Type >
        using Col_type = Map< uint, Cell< Cell_Type > * >
```

```
• template<typename Ta = double, typename Tb = uint> using MapVec_type = std::unordered_map< std::vector< Ta >, Tb, vecHasher< Ta >>
```

```
    template<typename Array_Type , typename Data_Type >
        using Counter_fun_type = std::function< double(const Array_Type &, uint, uint, Data_Type *)>
        Counter and rule functions.
```

```
    template<typename Array_Type , typename Data_Type >
        using Rule_fun_type = std::function< bool(const Array_Type &, uint, uint, Data_Type *)>
```

Functions

```
    template<typename T >
        bool vec_equal (const std::vector< T > &a, const std::vector< T > &b)
    Compares if -a- and -b- are equal.
```

```
• template<typename T > bool vec_equal_approx (const std::vector< T > &a, const std::vector< T > &b, double eps=1e-10)
```

Variables

```
const int CHECK::BOTH = -1
const int CHECK::NONE = 0
const int CHECK::ONE = 1
const int CHECK::TWO = 2
const int EXISTS::BOTH = -1
const int EXISTS::NONE = 0
const int EXISTS::ONE = 1
const int EXISTS::TWO = 1
```

- const int EXISTS::UKNOWN = -1
- const int EXISTS::AS_ZERO = 0
- const int EXISTS::AS_ONE = 1

8.46.1 Typedef Documentation

8.46.1.1 Col type

```
template<typename Cell_Type >
using Col_type = Map< uint, Cell<Cell_Type>* >
```

Definition at line 51 of file typedefs.hpp.

8.46.1.2 Counter_fun_type

```
template<typename Array_Type , typename Data_Type >
using Counter_fun_type = std::function<double(const Array_Type &, uint, uint, Data_Type *)>
```

Counter and rule functions.

Parameters

Array_Type	a BArray
unit,uint	Focal cell
Data_Type	Data associated with the function, for example, id of the attribute in the Array.

Returns

```
Counter_fun_type a double (the change statistic)
Rule_fun_type a bool. True if the cell is blocked.
```

Definition at line 123 of file typedefs.hpp.

8.46.1.3 Counts_type

```
typedef std::vector< std::pair< std::vector<double>, uint > > Counts_type
```

Definition at line 44 of file typedefs.hpp.

8.46.1.4 MapVec_type

```
template<typename Ta = double, typename Tb = uint>
using MapVec_type = std::unordered_map< std::vector< Ta >, Tb, vecHasher<Ta> >
```

Definition at line 105 of file typedefs.hpp.

8.46.1.5 Row_type

```
template<typename Cell_Type >
using Row_type = Map< uint, Cell<Cell_Type> >
```

Definition at line 48 of file typedefs.hpp.

8.46.1.6 Rule_fun_type

```
template<typename Array_Type , typename Data_Type >
using Rule_fun_type = std::function<bool(const Array_Type &, uint, uint, Data_Type *)>
```

Definition at line 126 of file typedefs.hpp.

8.46.1.7 uint

```
typedef unsigned int uint
```

Definition at line 10 of file typedefs.hpp.

8.46.2 Function Documentation

8.46.2.1 vec_equal()

Compares if -a- and -b- are equal.

Parameters

```
a,b Two vectors of the same length
```

Returns

true if all elements are equal.

Definition at line 137 of file typedefs.hpp.

8.46.2.2 vec_equal_approx()

Definition at line 155 of file typedefs.hpp.

8.46.2.3 vec_inner_prod()

Definition at line 175 of file typedefs.hpp.

8.47 README.md File Reference

Index

```
\simBArray
                                                     \simRules
                                                          Rules < Array_Type, Data_Type >, 152
    BArray< Cell Type, Data Type >, 31
\simBArrayCell
                                                     \simStatsCounter
    BArrayCell< Cell_Type, Data_Type >, 42
                                                          StatsCounter< Array_Type, Data_Type >, 156
~BArrayCell const
                                                     \simSupport
    BArrayCell_const< Cell_Type, Data_Type >, 44
                                                         Support<
                                                                      Array_Type,
                                                                                     Data Counter Type,
{\sim}\mathsf{BArrayDense}
                                                              Data_Rule_Type, Data_Rule_Dyn_Type >,
    BArrayDense < Cell_Type, Data_Type >, 50
\simBArrayDenseCell
                                                     add
    BArrayDenseCell< Cell_Type, Data_Type >, 62
                                                         Cell< Cell_Type >, 76, 77
~BArrayDenseCell const
                                                         FreqTable < T >, 100
    BArrayDenseCell const< Cell Type, Data Type
                                                     add array
         >, 65
                                                         Model<
                                                                                     Data Counter Type,
                                                                     Array_Type,
\simBArrayVector
                                                              Data_Rule_Type, Data_Rule_Dyn_Type >,
    BArrayVector< Cell_Type, Data_Type >, 68
~BArrayVector const
                                                     add counter
    BArrayVector_const< Cell_Type, Data_Type >, 71
                                                         Counters < Array_Type, Data_Type >, 88
\simCell
                                                         Model<
                                                                     Array Type,
                                                                                     Data Counter Type,
    Cell< Cell_Type >, 75
                                                              Data_Rule_Type, Data_Rule_Dyn_Type >,
\simConstBArrayRowIter
                                                              118
    ConstBArrayRowlter< Cell Type, Data Type >, 80
                                                          StatsCounter < Array_Type, Data_Type >, 156
\simCounter
                                                                      Array Type,
                                                                                     Data Counter Type,
                                                         Support<
    Counter< Array_Type, Data_Type >, 83
                                                              Data_Rule_Type, Data_Rule_Dyn_Type >,
\simCounters
    Counters < Array_Type, Data_Type >, 87
                                                     add data
\simEntries
                                                          Flock, 94
    Entries < Cell_Type >, 91
                                                     add_rule
\simFlock
                                                         Model <
                                                                     Array_Type,
                                                                                     Data_Counter_Type,
    Flock, 94
                                                              Data_Rule_Type, Data_Rule_Dyn_Type >,
\simFreqTable
    FreqTable < T >, 99
                                                         PowerSet< Array_Type, Data_Rule_Type >, 145
\simGeese
                                                         Rules < Array_Type, Data_Type >, 153
    Geese, 105
                                                         Support<
                                                                      Array Type,
                                                                                     Data Counter Type,
\simModel
                                                              Data_Rule_Type, Data_Rule_Dyn_Type >,
    Model<
                Array Type,
                                Data Counter Type,
                                                              162
         add rule dyn
         117
                                                         Model <
                                                                     Array_Type,
                                                                                     Data Counter Type,
\simNetCounterData
                                                              Data_Rule_Type, Data_Rule_Dyn_Type >,
    NetCounterData, 128
                                                              119, 120
\simNetworkData
                                                          Support<
                                                                      Array_Type,
                                                                                     Data_Counter_Type,
    NetworkData, 130
                                                              Data_Rule_Type, Data_Rule_Dyn_Type >,
\simNode
                                                              163
    Node, 133
                                                     annotations
\simPhyloRuleDynData
                                                         Node, 134
    PhyloRuleDynData, 141
                                                     Array
\simPowerSet
                                                          ConstBArrayRowIter< Cell Type, Data Type >, 80
    PowerSet < Array_Type, Data_Rule_Type >, 145
                                                     arrav
\simRule
                                                         Node, 134
    Rule < Array_Type, Data_Type >, 150
                                                     array_bank
```

support-meat.hpp, 235	BARRY_BARRAY_MEAT_OPERATORS_HPP, 174
arrays	
Node, 134	checkdim_, 174
AS_ONE	COL, 174
EXISTS, 25	ROW, 174
as_vector	barray-meat.hpp
FreqTable $<$ T $>$, 100	COL, 176
AS_ZERO	ROW, 176
EXISTS, 25	BARRAY_BONES_HPP
at	barray-bones.hpp, 172
PhyloCounterData, 139	BArrayCell
•	BArrayCell< Cell_Type, Data_Type >, 41
BArray	BArrayCell< Cell_Type, Data_Type >, 41
BArray< Cell_Type, Data_Type >, 30, 31	~BArrayCell, 42
BArray< Cell_Type, Data_Type >, 27	BArray< Cell_Type, Data_Type >, 40
\sim BArray, 31	BArrayCell, 41
BArray, 30, 31	BArrayDense< Cell_Type, Data_Type >, 61
BArrayCell< Cell_Type, Data_Type >, 40	operator Cell_Type, 42
BArrayCell_const< Cell_Type, Data_Type >, 40	
clear, 31	operator*=, 42
col, 31	operator+=, 42
D, 32	operator-=, 42
default_val, 32	operator/=, 43
	operator=, 43
flush_data, 32	operator==, 43
get_cell, 32	BArrayCell_const
get_col_vec, 32, 33	BArrayCell_const< Cell_Type, Data_Type >, 44
get_entries, 33	BArrayCell_const< Cell_Type, Data_Type >, 43
get_row_vec, 33	\sim BArrayCell_const, 44
insert_cell, 33, 34	BArray Cell_Type, Data_Type >, 40
is_empty, 34	BArrayCell_const, 44
ncol, 34	BArrayDense < Cell_Type, Data_Type >, 61
nnozero, 34	operator Cell_Type, 44
nrow, 35	operator!=, 45
operator*=, 35	operator<, 45
operator(), 35	operator $<$, 45
operator+=, 35, 36	•
operator-=, 36	operator>, 45
operator/=, 36	operator>=, 46
operator=, 36, 37	operator==, 45
operator==, 37	BArrayDense
out_of_range, 37	BArrayDense< Cell_Type, Data_Type >, 49, 50
print, 37	BArrayDense < Cell_Type, Data_Type >, 46
•	\sim BArrayDense, 50
reserve, 37	BArrayCell< Cell_Type, Data_Type >, 61
resize, 37	BArrayCell_const< Cell_Type, Data_Type >, 61
rm_cell, 38	BArrayDense, 49, 50
row, 38	clear, 50
set_data, 38	col, <u>51</u>
swap_cells, 38	D, 51
swap_cols, 39	default_val, 51
swap_rows, 39	get_cell, 51
toggle_cell, 39	get_col_vec, 52
toggle_lock, 39	get_entries, 52
transpose, 39	get_row_vec, 52, 53
visited, 40	insert_cell, 53
zero_col, 40	
zero_row, 40	is_empty, 54
barray-bones.hpp	ncol, 54
BARRAY_BONES_HPP, 172	nnozero, 54
barray-meat-operators.hpp	nrow, 54

operator*=, 55	BArrayVector< Cell_Type, Data_Type >, 67
operator(), 54, 55	BArrayVector< Cell_Type, Data_Type >, 67
operator+=, 55	\sim BArrayVector, 68
operator-=, 56	BArrayVector, 67
operator/=, 56	begin, 68
operator=, 56, 57	end, 68
operator==, 57	is_col, 68
out_of_range, 57	is_row, 69
print, 57	operator std::vector< Cell_Type >, 69
reserve, 57	operator*=, 69
resize, 58	operator+=, 69
rm_cell, 58	operator-=, 69
row, 58	operator/=, 70
set_data, 58	operator=, 70
swap_cells, 59	operator==, 70
swap_cols, 59	size, 70
swap_rows, 59	barrayvector-meat.hpp
toggle_cell, 59	BARRY_BARRAYVECTOR_MEAT_HPP, 185
toggle_lock, 60	BArrayVector_const
transpose, 60	BArrayVector_const< Cell_Type, Data_Type >, 71
visited, 61	BArrayVector_const< Cell_Type, Data_Type >, 71
zero_col, 60	~BArrayVector_const, 71
zero_row, 60	BArrayVector_const, 71
barraydense-meet.hpp	begin, 72
BARRY_BARRAYDENSE_MEAT_HPP, 180	end, 72
COL, 180	is_col, 72
POS, 181	is_row, 72
ROW, 181	operator std::vector< Cell_Type >, 72
ZERO_CELL, 181	operator!=, 72
BArrayDenseCell	operator<, 73
BArrayDenseCell< Cell_Type, Data_Type >, 62	operator<=, 73
BArrayDenseCell< Cell_Type, Data_Type >, 61	operator>, 73
~BArrayDenseCell, 62	operator>=, 73
BArrayDenseCell, 62	operator==, 73
operator Cell_Type, 62	size, 74
operator*=, 63	barry, 23
operator+=, 63	barry-configuration.hpp
operator-=, 63	BARRY_CHECK_SUPPORT, 186
operator/=, 63	BARRY ISFINITE, 186
operator=, 63	BARRY_MAX_NUM_ELEMENTS, 186
operator==, 64	BARRY_SAFE_EXP, 186
barraydensecell-meat.hpp	Map, 186
BARRY BARRAYDENSECELL MEAT HPP, 183	printf_barry, 186
POS, 183	barry.hpp
BArrayDenseCell_const	BARRY_HPP, 188
BArrayDenseCell_const< Cell_Type, Data_Type	BARRY_VERSION, 188
	COUNTER FUNCTION, 188
>, 65 PArroy Depos Cell const < Cell Type Data Type > 64	COUNTER_LAMBDA, 188
BArrayDenseCell_const< Cell_Type, Data_Type >, 64	RULE FUNCTION, 189
~BArrayDenseCell_const, 65 BArrayDenseCell_const, 65	
	RULE_LAMBDA, 189
operator Cell_Type, 65	barry::counters, 23
operator!=, 65	barry::counters::network, 24
operator<, 65	barry::counters::phylo, 24
operator<=, 66	BARRY_BARRAY_MEAT_OPERATORS_HPP
operator>, 66	barray-meat-operators.hpp, 174
operator>=, 66	BARRY_BARRAYDENSE_MEAT_HPP
operator==, 66	barraydense-meet.hpp, 180
BArrayVector	BARRY_BARRAYDENSECELL_MEAT_HPP

harrandanaaall maat han 100	NONE 24
barraydensecell-meat.hpp, 183 BARRY BARRAYVECTOR MEAT HPP	NONE, 24 ONE, 24
barrayvector-meat.hpp, 185	TWO, 24
BARRY_CHECK_SUPPORT	checkdim
barry-configuration.hpp, 186	barray-meat-operators.hpp, 174
BARRY HPP	clear
barry.hpp, 188	BArray< Cell_Type, Data_Type >, 31
BARRY_ISFINITE	BArrayDense< Cell_Type, Data_Type >, 50
barry-configuration.hpp, 186	Counters < Array_Type, Data_Type >, 88
BARRY_MAX_NUM_ELEMENTS	FreqTable < T >, 100
barry-configuration.hpp, 186	Rules< Array_Type, Data_Type >, 153
BARRY_SAFE_EXP	COL
barry-configuration.hpp, 186	barray-meat-operators.hpp, 174
BARRY_SUPPORT_MEAT_HPP	barray-meat.hpp, 176
support-meat.hpp, 229	barraydense-meet.hpp, 180
BARRY_VERSION	col
barry.hpp, 188	BArray< Cell_Type, Data_Type >, 31
begin	BArrayDense < Cell_Type, Data_Type >, 51
BArrayVector < Cell_Type, Data_Type >, 68	Col_type
BArrayVector_const< Cell_Type, Data_Type >, 72	typedefs.hpp, 240
PhyloCounterData, 139 PowerSet< Array_Type, Data_Rule_Type >, 145	colnames Flock, 94
blengths	Geese, 105
NodeData, 137	Model < Array_Type, Data_Counter_Type,
BOTH	Data_Rule_Type, Data_Rule_Dyn_Type >,
CHECK, 24	120
EXISTS, 25	conditional_prob
, -	Model< Array_Type, Data_Counter_Type,
calc	Data_Rule_Type, Data_Rule_Dyn_Type >,
PowerSet < Array_Type, Data_Rule_Type >, 146	120
Support< Array_Type, Data_Counter_Type,	ConstBArrayRowIter
Data_Rule_Type, Data_Rule_Dyn_Type >,	ConstBArrayRowIter< Cell_Type, Data_Type >, 79
163	ConstBArrayRowIter< Cell_Type, Data_Type >, 79
calc_backend	\sim ConstBArrayRowIter, 80
support-meat.hpp, 230	Array, 80
calc_reduced_sequence	ConstBArrayRowIter, 79
Geese, 105 calc_sequence	current_col, 80
Geese, 105	current_row, 80
Cell	iter, 80
Cell< Cell_Type >, 75, 76	coordinates_free
Cell< Cell_Type >, 74	PowerSet < Array_Type, Data_Rule_Type >, 147 Support < Array_Type, Data_Counter_Type,
~Cell, 75	Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,
add, 76, 77	167
Cell, 75, 76	coordinates_locked
operator Cell_Type, 77	PowerSet < Array_Type, Data_Rule_Type >, 147
operator!=, 77	Support< Array_Type, Data_Counter_Type,
operator=, 77, 78	Data_Rule_Type, Data_Rule_Dyn_Type >,
operator==, 78	167
value, 78	count
visited, 78	Counter< Array_Type, Data_Type >, 83
cfree	count_all
support-meat.hpp, 235	StatsCounter< Array_Type, Data_Type >, 157
change_stats	count_current
Support	StatsCounter< Array_Type, Data_Type >, 157
Data_Rule_Type, Data_Rule_Dyn_Type >,	count_fun
166	Counter< Array_Type, Data_Type >, 84
CHECK, 24	count_init
BOTH, 24	

StatsCounter< Array_Type, Data_Type >, 157	Phylo counters, 18
Counter	counter_mutual
Counter< Array_Type, Data_Type >, 82, 83	Network counters, 13
Counter< Array_Type, Data_Type >, 81	counter_neofun
\sim Counter, 83	Phylo counters, 19
count, 83	counter_neofun_a2b
count_fun, 84	Phylo counters, 19
Counter, 82, 83	counter_nodecov
data, 84	Network counters, 13
delete_data, 85	counter_nodeicov
desc, 85	Network counters, 14
init, 83	counter_nodematch
init_fun, 85	Network counters, 14
name, 85	counter_nodeocov
operator=, 84	Network counters, 14
counter_absdiff	counter_odegree
Network counters, 11	Network counters, 14
counter_co_opt	counter_odegree15
Phylo counters, 16	Network counters, 14
counter_cogain	counter_ostar2
Phylo counters, 17	Network counters, 15
counter_css_partially_false_recip	counter_overall_changes
Network counters, 11	Phylo counters, 19
counter_ctriads	counter_overall_gains
Network counters, 11	Phylo counters, 19
counter_degree	counter_overall_loss
Network counters, 11	Phylo counters, 20
counter_density	counter_prop_genes_changing
Network counters, 12	Phylo counters, 20
counter_diff	counter_subfun
Network counters, 12	Phylo counters, 20
counter_edges	counter_ttriads
Network counters, 12	Network counters, 15
Counter_fun_type	Counters
typedefs.hpp, 240	Counters< Array_Type, Data_Type >, 86, 87
COUNTER_FUNCTION	counters
barry.hpp, 188	support-meat.hpp, 235
counter_gains	Counters< Array_Type, Data_Type >, 86
Phylo counters, 17	~Counters, 87
counter_gains_k_offspring	add counter, 88
Phylo counters, 17	clear, 88
counter_genes_changing	Counters, 86, 87
Phylo counters, 18	operator=, 88, 89
counter idegree	operator[], 89
Network counters, 12	size, 90
counter_idegree15	
	counters_
Network counters, 13	support-meat.hpp, 235
counter_isolates	Counting, 9
Network counters, 13	Counts
counter_istar2	PhyloRuleDynData, 142
Network counters, 13	Counts_type
COUNTER_LAMBDA	typedefs.hpp, 240
barry.hpp, 188	current_col
counter_longest	ConstBArrayRowIter< Cell_Type, Data_Type >, 80
Phylo counters, 18	current_row
counter_loss	ConstBArrayRowIter< Cell_Type, Data_Type >, 80
Phylo counters, 18	current_stats
counter_maxfuns	Support< Array_Type, Data_Counter_Type,

```
\simEntries, 91
         Data_Rule_Type, Data_Rule_Dyn_Type >,
                                                           Entries, 91
                                                           resize, 91
D
                                                           source, 92
    BArray< Cell_Type, Data_Type >, 32
                                                           target, 92
    BArrayDense < Cell Type, Data Type >, 51
                                                           val, 92
    Rule < Array_Type, Data_Type >, 151
                                                      eval rules dyn
dat
                                                           Support<
                                                                                       Data Counter Type,
                                                                        Array Type,
    Flock, 98
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
data
    Counter< Array_Type, Data_Type >, 84
                                                      EXISTS, 25
    PowerSet < Array_Type, Data_Rule_Type >, 148
                                                           AS ONE, 25
default val
                                                           AS ZERO, 25
    BArray< Cell Type, Data Type >, 32
                                                           BOTH, 25
    BArrayDense < Cell_Type, Data_Type >, 51
                                                           NONE, 26
delete counters
                                                           ONE, 26
    Support<
                 Array_Type,
                                 Data_Counter_Type,
                                                           TWO, 26
         Data_Rule_Type, Data_Rule_Dyn_Type >,
                                                           UKNOWN, 26
    support-meat.hpp, 236
                                                           support-meat.hpp, 236
delete data
                                                      Flock, 92
    Counter< Array Type, Data Type >, 85
                                                           \simFlock, 94
delete rengine
                                                           add data, 94
    Geese, 112
                                                           colnames, 94
delete rules
                                                           dat, 98
    Support<
                 Array Type,
                                 Data Counter Type,
                                                           Flock, 93
         Data_Rule_Type, Data_Rule_Dyn_Type >,
                                                           get counters, 94
                                                           get_model, 95
    support-meat.hpp, 236
                                                           get_support, 95
delete_rules_dyn
                                                           init, 95
                                 Data Counter Type,
    Support<
                 Array Type,
                                                           initialized, 98
         Data_Rule_Type, Data_Rule_Dyn_Type >,
                                                           likelihood joint, 95
         168
                                                           model, 98
    support-meat.hpp, 236
                                                           nfunctions, 98
delete_support
                                                           nfuns. 96
    Geese, 112
                                                           nleafs, 96
desc
    Counter< Array_Type, Data_Type >, 85
                                                           nnodes, 96
                                                           nterms, 96
directed
                                                           ntrees, 96
    NetworkData, 130
                                                           operator(), 96
duplication
                                                           parse polytomies, 97
    Node, 135
                                                           print. 97
    NodeData, 138
                                                           rengine, 98
    PhyloRuleDynData, 142
                                                           set seed, 97
else
                                                           support size, 97
    support-meat.hpp, 236
                                                      flush data
empty
                                                           BArray< Cell_Type, Data_Type >, 32
    PhyloCounterData, 139
                                                      for
EmptyArray
                                                           support-meat.hpp, 231
    PowerSet < Array_Type, Data_Rule_Type >, 148
                                                      FreqTable
end
                                                           FreqTable < T >, 99
    BArrayVector< Cell Type, Data Type >, 68
                                                      FregTable < T >, 99
    BArrayVector_const< Cell_Type, Data_Type >, 72
                                                           \simFregTable, 99
    PhyloCounterData, 139
                                                           add. 100
    PowerSet< Array Type, Data Rule Type >, 146
                                                           as_vector, 100
Entries
                                                           clear, 100
    Entries < Cell_Type >, 91
                                                           FreqTable, 99
Entries < Cell_Type >, 90
                                                           get_data, 100
```

get_data_ptr, 100 print, 100 reserve, 101 size, 101	Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 121 get_annotated_nodes
, -	Geese, 105
Geese, 101	get_cell
\sim Geese, 105	BArray< Cell_Type, Data_Type >, 32
calc_reduced_sequence, 105	BArrayDense< Cell_Type, Data_Type >, 51
calc_sequence, 105	get_col_vec
colnames, 105	BArray< Cell_Type, Data_Type >, 32, 33
delete_rengine, 112	BArrayDense< Cell Type, Data Type >, 52
delete_support, 112	get_counters
Geese, 104	Flock, 94
get_annotated_nodes, 105	Geese, 106
get_counters, 106	Model Array_Type, Data_Counter_Type,
get_model, 106	Data_Rule_Type, Data_Rule_Dyn_Type >,
get_probabilities, 106	121
get_rengine, 106	PhyloCounterData, 140
get_states, 106	StatsCounter< Array_Type, Data_Type >, 157
get_support, 107	Support< Array_Type, Data_Counter_Type,
inherit_support, 107	Data_Rule_Type, Data_Rule_Dyn_Type >,
init, 107	164
init_node, 107	get_counts
initialized, 112	Support< Array_Type, Data_Counter_Type,
likelihood, 107	Data_Rule_Type, Data_Rule_Dyn_Type >,
likelihood_exhaust, 108	164
map_to_nodes, 112	get_counts_ptr
nannotations, 108	Support< Array_Type, Data_Counter_Type,
nfunctions, 112	Data_Rule_Type, Data_Rule_Dyn_Type >,
nfuns, 108	164
nleafs, 108	get_current_stats
nnodes, 108	Support< Array_Type, Data_Counter_Type,
nodes, 113	Data_Rule_Type, Data_Rule_Dyn_Type >,
nterms, 109	164
observed_counts, 109	get_data
operator=, 109	FreqTable $<$ T $>$, 100
parse_polytomies, 109	PowerSet < Array_Type, Data_Rule_Type >, 146
predict, 109	Support< Array_Type, Data_Title_Type >, Table Support< Array_Type, Data_Counter_Type,
predict_backend, 110	Data_Rule_Type, Data_Rule_Dyn_Type >,
predict_exhaust, 110	164
predict_exhaust_backend, 110	get data ptr
predict_sim, 110	FreqTable < T >, 100
print, 110	PowerSet< Array Type, Data Rule Type >, 146
print_observed_counts, 111	get entries
reduced sequence, 113	BArray< Cell_Type, Data_Type >, 33
sequence, 113	BArrayDense < Cell Type, Data Type >, 52
set_seed, 111	get last name
simulate, 111	phylo.hpp, 205
support_size, 111	get_model
update_annotations, 111	Flock, 95
geese-bones.hpp	Geese, 106
INITIALIZED, 211	get_norm_const
keygen_full, 212	Model< Array_Type, Data_Counter_Type,
RULE_FUNCTION, 212	Data_Rule_Type, Data_Rule_Dyn_Type >,
vec_diff, 212	121
vector_caster, 212	get parent
gen key	Node, 133
9·,	get_probabilities
	ger_hi onaniillies

Geese, 106	include/barry/barry.hpp, 187
get_pset	include/barry/cell-bones.hpp, 189
Model< Array_Type, Data_Counter_Type,	include/barry/cell-meat.hpp, 190
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/col-bones.hpp, 191
121	include/barry/counters-bones.hpp, 191
get_pset_stats	include/barry/counters-meat.hpp, 192
Model< Array_Type, Data_Counter_Type,	include/barry/counters/network.hpp, 193
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/counters/phylo.hpp, 200
121	include/barry/model-bones.hpp, 205
get_rengine	include/barry/model-meat.hpp, 207
Geese, 106	include/barry/models/geese.hpp, 209
Model < Array_Type, Data_Counter_Type,	include/barry/models/geese/flock-bones.hpp, 210
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/models/geese/flock-meet.hpp, 210
122	include/barry/models/geese/geese-bones.hpp, 211
get_row_vec	include/barry/models/geese/geese-meat-constructors.hpp,
BArray< Cell_Type, Data_Type >, 33	213
BArrayDense< Cell_Type, Data_Type >, 52, 53	include/barry/models/geese/geese-meat-likelihood.hpp,
get_rules	213
Model< Array_Type, Data_Counter_Type,	include/barry/models/geese/geese-meat-likelihood_exhaust.hpp,
Data_Rule_Type, Data_Rule_Dyn_Type >,	214
122	include/barry/models/geese/geese-meat-predict.hpp,
Support< Array_Type, Data_Counter_Type,	215
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/models/geese/geese-meat-predict_exhaust.hpp,
165	215
get_rules_dyn	include/barry/models/geese/geese-meat-predict_sim.hpp,
Model Array_Type, Data_Counter_Type,	216
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/models/geese/geese-meat-simulate.hpp,
122	216
Support< Array_Type, Data_Counter_Type,	include/barry/models/geese/geese-meat.hpp, 217
Data_Rule_Type, Data_Bule_Dyn_Type >,	include/barry/models/geese/geese-node-bones.hpp,
165	217
	include/barry/powerset-bones.hpp, 218
get_seq Pulos < Array Type Data Type > 152	include/barry/powerset-bories.hpp, 219
Rules < Array_Type, Data_Type >, 153	include/barry/rules-bones.hpp, 221
get_states	include/barry/rules-bories.hpp, 221
Geese, 106	include/barry/statscounter-bones.hpp, 223
get_support	•
Flock, 95	include/barry/statscounter-meat.hpp, 224
Geese, 107	include/barry/statsdb.hpp, 225
Model Array_Type, Data_Counter_Type,	include/barry/support-bones.hpp, 226
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/support-meat.hpp, 228
122	include/barry/typedefs.hpp, 238
id	indices
Node, 135	NetCounterData, 128
if	inherit_support
support-meat.hpp, 231	Geese, 107
include/barry/barray-bones.hpp, 171	init
include/barry/barray-iterator.hpp, 172	Counter< Array_Type, Data_Type >, 83
include/barry/barray-meat-operators.hpp, 173	Flock, 95
include/barry/barray-meat-operators.npp, 175	Geese, 107
• • • • • • • • • • • • • • • • • • • •	init_fun
include/barry/barraycell-bones.hpp, 176	Counter< Array_Type, Data_Type >, 85
include/barry/barraydense hance han 179	init_node
include/barry/barraydense-bones.hpp, 178	Geese, 107
include/barry/barraydense-meet.hpp, 179	init_support
include/barry/barraydensecell-bones.hpp, 181	PowerSet < Array_Type, Data_Rule_Type >, 146
include/barry/barraydensecell-meat.hpp, 182	Support< Array_Type, Data_Counter_Type,
include/barry/barrayvector-bones.hpp, 183	Data_Rule_Type, Data_Rule_Dyn_Type >,
include/barry/barrayvector-meat.hpp, 184	165
include/barry/barry-configuration.hpp, 185	

INITIALIZED	Support< Array_Type, Data_Counter_Type,
geese-bones.hpp, 211	Data_Rule_Type, Data_Rule_Dyn_Type >,
initialized	168
Flock, 98	Model
Geese, 112	Model< Array_Type, Data_Counter_Type,
insert_cell	$Data_Rule_Type, Data_Rule_Dyn_Type >,$
BArray< Cell_Type, Data_Type >, 33, 34	117
BArrayDense< Cell_Type, Data_Type >, 53	model
support-meat.hpp, 231	Flock, 98
is_col	Model < Array_Type, Data_Counter_Type, Data_Rule_Type
BArrayVector< Cell_Type, Data_Type >, 68	Data_Rule_Dyn_Type >, 113
BArrayVector_const< Cell_Type, Data_Type >, 72	\sim Model, 117
is_empty	add_array, 118
BArray< Cell_Type, Data_Type >, 34	add_counter, 118
BArrayDense< Cell_Type, Data_Type >, 54	add_rule, 119
is_leaf	add_rule_dyn, 119, 120
Node, 134	colnames, 120
is_row	conditional_prob, 120
BArrayVector< Cell_Type, Data_Type >, 69	gen_key, 121
BArrayVector_const< Cell_Type, Data_Type >, 72	get_counters, 121
iter	get_norm_const, 121
ConstBArrayRowIter< Cell_Type, Data_Type >, 80	get_pset, 121
	get_pset_stats, 121
keygen_default	get_rengine, 122
model-bones.hpp, 206	get_rules, 122
keygen_full	get_rules_dyn, 122
geese-bones.hpp, 212	get_support, 122
lla.	likelihood, 122, 123
Bhyla Byla Dya Data 140	likelihood_total, 123
PhyloRuleDynData, 142	Model, 117
likelihood	nterms, 123
Geese, 107	operator=, 124
Model Array_Type, Data_Counter_Type,	print, 124
Data_Rule_Type, Data_Rule_Dyn_Type >, 122, 123	print_stats, 124
likelihood_	sample, 124
model-meat.hpp, 208	set_counters, 125
• •	set_keygen, 125
likelihood_exhaust Geese, 108	set_rengine, 125
likelihood_joint	set_rules, 125
Flock, 95	set_rules_dyn, 125
likelihood_total	set_seed, 126
Model Array_Type, Data_Counter_Type,	size, 126
Data Rule Type, Data Rule Dyn Type >,	size_unique, 126
123	store_psets, 126
123	support_size, 126
M	model-bones.hpp
PowerSet < Array_Type, Data_Rule_Type >, 148	keygen_default, 206
Support< Array_Type, Data_Counter_Type,	model-meat.hpp
Data_Rule_Type, Data_Rule_Dyn_Type >,	likelihood_, 208
168	MODEL_TEMPLATE, 207, 208
Мар	MODEL_TEMPLATE_ARGS, 207
barry-configuration.hpp, 186	MODEL_TYPE, 208
map_to_nodes	update_normalizing_constant, 209
Geese, 112	MODEL_TEMPLATE
MapVec_type	model-meat.hpp, 207, 208
typedefs.hpp, 240	MODEL_TEMPLATE_ARGS
max_num_elements	model-meat.hpp, 207
	MODEL_TYPE

mandal manak hasa 000	animates adams of 5.4.4
model-meat.hpp, 208	counter_odegree15, 14
N	counter_ostar2, 15
	counter_ttriads, 15
PowerSet< Array_Type, Data_Rule_Type >, 148 Support< Array_Type, Data_Counter_Type,	NETWORK_COUNTER, 15
= =	network.hpp
Data_Rule_Type, Data_Rule_Dyn_Type >,	NET_C_DATA_IDX, 196
168	NET_C_DATA_NUM, 196
name	NetCounter, 198
Counter< Array_Type, Data_Type >, 85	NetCounters, 198
nannotations	NetModel, 198
Geese, 108	NetRule, 198
narray	NetRules, 198
Node, 135	NetStatsCounter, 198
ncol	NetSupport, 199
BArray< Cell_Type, Data_Type >, 34	• •
BArrayDense< Cell_Type, Data_Type >, 54	Network, 199
NET C DATA IDX	NETWORK_COUNTER, 196
	NETWORK_COUNTER_LAMBDA, 197
network.hpp, 196	NETWORK_RULE, 197
NET_C_DATA_NUM	NETWORK_RULE_LAMBDA, 197
network.hpp, 196	rules_zerodiag, 199
NetCounter	NETWORK COUNTER
network.hpp, 198	Network counters, 15
NetCounterData, 127	network.hpp, 196
∼NetCounterData, 128	NETWORK_COUNTER_LAMBDA
indices, 128	
NetCounterData, 127	network.hpp, 197
numbers, 128	NETWORK_RULE
NetCounters	network.hpp, 197
	NETWORK_RULE_LAMBDA
network.hpp, 198	network.hpp, 197
NetModel	NetworkData, 128
network.hpp, 198	\sim NetworkData, 130
NetRule	directed, 130
network.hpp, 198	NetworkData, 129, 130
NetRules	vertex_attr, 130
network.hpp, 198	nfunctions
NetStatsCounter	Flock, 98
network.hpp, 198	
NetSupport	Geese, 112
network.hpp, 199	nfuns
Network	Flock, 96
	Geese, 108
network.hpp, 199	nleafs
Network counters, 10	Flock, 96
counter_absdiff, 11	Geese, 108
counter_css_partially_false_recip, 11	nnodes
counter_ctriads, 11	
counter_degree, 11	Flock, 96
counter density 10	•
counter_density, 12	Geese, 108
_ ·	Geese, 108 nnozero
counter_diff, 12	Geese, 108 nnozero BArray< Cell_Type, Data_Type >, 34
counter_diff, 12 counter_edges, 12	Geese, 108 nnozero BArray< Cell_Type, Data_Type >, 34 BArrayDense< Cell_Type, Data_Type >, 54
counter_diff, 12 counter_edges, 12 counter_idegree, 12	Geese, 108 nnozero BArray< Cell_Type, Data_Type >, 34 BArrayDense< Cell_Type, Data_Type >, 54 Node, 131
counter_diff, 12 counter_edges, 12 counter_idegree, 12 counter_idegree15, 13	Geese, 108 nnozero BArray< Cell_Type, Data_Type >, 34 BArrayDense< Cell_Type, Data_Type >, 54 Node, 131 ~Node, 133
counter_diff, 12 counter_edges, 12 counter_idegree, 12 counter_idegree15, 13 counter_isolates, 13	Geese, 108 nnozero BArray< Cell_Type, Data_Type >, 34 BArrayDense< Cell_Type, Data_Type >, 54 Node, 131 ~Node, 133 annotations, 134
counter_diff, 12 counter_edges, 12 counter_idegree, 12 counter_idegree15, 13 counter_isolates, 13 counter_istar2, 13	Geese, 108 nnozero BArray< Cell_Type, Data_Type >, 34 BArrayDense< Cell_Type, Data_Type >, 54 Node, 131 ~Node, 133
counter_diff, 12 counter_edges, 12 counter_idegree, 12 counter_idegree15, 13 counter_isolates, 13 counter_istar2, 13 counter_mutual, 13	Geese, 108 nnozero BArray< Cell_Type, Data_Type >, 34 BArrayDense< Cell_Type, Data_Type >, 54 Node, 131 ~Node, 133 annotations, 134
counter_diff, 12 counter_edges, 12 counter_idegree, 12 counter_idegree15, 13 counter_isolates, 13 counter_istar2, 13 counter_mutual, 13 counter_nodecov, 13	Geese, 108 nnozero BArray< Cell_Type, Data_Type >, 34 BArrayDense< Cell_Type, Data_Type >, 54 Node, 131 ~Node, 133 annotations, 134 array, 134
counter_diff, 12 counter_edges, 12 counter_idegree, 12 counter_idegree15, 13 counter_isolates, 13 counter_istar2, 13 counter_mutual, 13	Geese, 108 nnozero BArray< Cell_Type, Data_Type >, 34 BArrayDense< Cell_Type, Data_Type >, 54 Node, 131 ~Node, 133 annotations, 134 array, 134 arrays, 134 duplication, 135
counter_diff, 12 counter_edges, 12 counter_idegree, 12 counter_idegree15, 13 counter_isolates, 13 counter_istar2, 13 counter_mutual, 13 counter_nodecov, 13	Geese, 108 nnozero BArray< Cell_Type, Data_Type >, 34 BArrayDense< Cell_Type, Data_Type >, 54 Node, 131 ~Node, 133 annotations, 134 array, 134 arrays, 134 duplication, 135 get_parent, 133
counter_diff, 12 counter_edges, 12 counter_idegree, 12 counter_idegree15, 13 counter_isolates, 13 counter_istar2, 13 counter_mutual, 13 counter_nodecov, 13 counter_nodeicov, 14	Geese, 108 nnozero BArray< Cell_Type, Data_Type >, 34 BArrayDense< Cell_Type, Data_Type >, 54 Node, 131 ~Node, 133 annotations, 134 array, 134 arrays, 134 duplication, 135

narray, 135	BArrayCell_const< Cell_Type, Data_Type >, 45
Node, 132, 133	BArrayDenseCell_const< Cell_Type, Data_Type
noffspring, 134	>, 65
offspring, 135	BArrayVector_const< Cell_Type, Data_Type >, 73
ord, 135	operator<=
parent, 136	BArrayCell_const< Cell_Type, Data_Type >, 45
probability, 136	BArrayDenseCell_const< Cell_Type, Data_Type
subtree_prob, 136	>, 66
visited, 136	BArrayVector_const< Cell_Type, Data_Type >, 73
NodeData, 137	operator>
blengths, 137	BArrayCell_const< Cell_Type, Data_Type >, 45
duplication, 138	BArrayDenseCell_const< Cell_Type, Data_Type
NodeData, 137	>, 66
states, 138	BArrayVector_const< Cell_Type, Data_Type >, 73
nodes	operator>=
	BArrayCell_const< Cell_Type, Data_Type >, 46
Geese, 113	BArrayDenseCell_const< Cell_Type, Data_Type BArrayDenseCell_const< Cell_Type, Data_Type
noffspring	
Node, 134	>, 66
NONE	BArrayVector_const< Cell_Type, Data_Type >, 73
CHECK, 24	operator*=
EXISTS, 26	BArray< Cell_Type, Data_Type >, 35
nrow	BArrayCell< Cell_Type, Data_Type >, 42
BArray< Cell_Type, Data_Type >, 35	BArrayDense < Cell_Type, Data_Type >, 55
BArrayDense < Cell_Type, Data_Type >, 54	BArrayDenseCell< Cell_Type, Data_Type >, 63
nterms	BArrayVector< Cell_Type, Data_Type >, 69
Flock, 96	operator()
Geese, 109	BArray< Cell_Type, Data_Type >, 35
Model< Array_Type, Data_Counter_Type,	BArrayDense < Cell_Type, Data_Type >, 54, 55
Data_Rule_Type, Data_Rule_Dyn_Type >,	Flock, 96
123	PhyloCounterData, 140
ntrees	Rule < Array_Type, Data_Type >, 151
Flock, 96	Rules < Array_Type, Data_Type >, 154
numbers	vecHasher $<$ T $>$, 169
NetCounterData, 128	operator+=
	BArray< Cell_Type, Data_Type >, 35, 36
observed_counts	BArrayCell< Cell_Type, Data_Type >, 42
Geese, 109	BArrayDense < Cell_Type, Data_Type >, 55
offspring	BArrayDenseCell< Cell_Type, Data_Type >, 63
Node, 135	BArrayVector< Cell_Type, Data_Type >, 69
ONE	operator-=
CHECK, 24	BArray< Cell_Type, Data_Type >, 36
EXISTS, 26	BArrayCell< Cell Type, Data Type >, 42
operator Cell_Type	BArrayDense< Cell Type, Data Type >, 56
BArrayCell< Cell_Type, Data_Type >, 42	BArrayDenseCell< Cell_Type, Data_Type >, 63
BArrayCell_const< Cell_Type, Data_Type >, 44	BArrayVector< Cell_Type, Data_Type >, 69
BArrayDenseCell< Cell_Type, Data_Type >, 62	operator/=
BArrayDenseCell_const< Cell_Type, Data_Type	BArray< Cell_Type, Data_Type >, 36
>, 65	BArrayCell< Cell_Type, Data_Type >, 43
Cell< Cell_Type >, 77	BArrayDense< Cell_Type, Data_Type >, 56
operator std::vector< Cell_Type >	BArrayDenseCell< Cell_Type, Data_Type >, 63
BArrayVector< Cell_Type, Data_Type >, 69	BArray Vector < Cell_Type, Data_Type >, 70
BArrayVector_const< Cell_Type, Data_Type >, 72	operator=
operator!=	·
BArrayCell_const< Cell_Type, Data_Type >, 45	BArray Cell_Type, Data_Type >, 36, 37
BArrayDenseCell_const< Cell_Type, Data_Type	BArrayCell< Cell_Type, Data_Type >, 43 BArrayConso< Coll_Type_Data_Type > 56, 57
>, 65	BArrayDense< Cell_Type, Data_Type >, 56, 57
BArrayVector_const< Cell_Type, Data_Type >, 72	BArrayDenseCell< Cell_Type, Data_Type >, 63
Cell< Cell_Type >, 77	BArrayVector< Cell_Type, Data_Type >, 70
operator<	Cell< Cell_Type >, 77, 78
-p	

Counter< Array_Type, Data_Type >, 84	PhyloPowerSet, 203
Counters < Array_Type, Data_Type >, 88, 89	PhyloRule, 204
Geese, 109	PhyloRuleData, 204
Model< Array_Type, Data_Counter_Type,	PhyloRuleDyn, 204
Data_Rule_Type, Data_Rule_Dyn_Type >,	PhyloRules, 204
124	PhyloRulesDyn, 204
Rules < Array_Type, Data_Type >, 154	PhyloStatsCounter, 204
operator==	PhyloSupport, 205
BArray< Cell_Type, Data_Type >, 37	PHYLO_CHECK_MISSING
BArrayCell< Cell_Type, Data_Type >, 43	phylo.hpp, 202
BArrayCell_const< Cell_Type, Data_Type >, 45	PHYLO_COUNTER_LAMBDA
BArrayDense< Cell_Type, Data_Type >, 57	phylo.hpp, 202
BArrayDenseCell< Cell_Type, Data_Type >, 64	PHYLO_RULE_DYN_LAMBDA
BArrayDenseCell_const< Cell_Type, Data_Type	phylo.hpp, 202
>, 66	PhyloArray
BArrayVector< Cell_Type, Data_Type >, 70	phylo.hpp, 203
BArrayVector_const< Cell_Type, Data_Type >, 73	PhyloCounter
Cell< Cell_Type >, 78	phylo.hpp, 203
operator[]	PhyloCounterData, 138
Counters< Array_Type, Data_Type >, 89	at, 139
PowerSet< Array_Type, Data_Rule_Type >, 147	begin, 139
ord	empty, 139
Node, 135	end, 139
out_of_range	get_counters, 140
-	operator(), 140
BArray Cell_Type, Data_Type >, 37	* **
BArrayDense< Cell_Type, Data_Type >, 57	PhyloCounterData, 139
parent	push_back, 140
Node, 136	reserve, 140
parse_polytomies	shrink_to_fit, 140
Flock, 97	size, 140
Geese, 109	PhyloCounters
	phylo.hpp, 203
Phylo counters, 15	PhyloModel
counter_co_opt, 16	phylo.hpp, 203
counter_cogain, 17	PhyloPowerSet
counter_gains, 17	phylo.hpp, 203
counter_gains_k_offspring, 17	PhyloRule
counter_genes_changing, 18	phylo.hpp, 204
counter_longest, 18	PhyloRuleData
counter_loss, 18	phylo.hpp, 204
counter_maxfuns, 18	PhyloRuleDyn
counter_neofun, 19	phylo.hpp, 204
counter_neofun_a2b, 19	PhyloRuleDynData, 141
counter_overall_changes, 19	~PhyloRuleDynData, 141
counter_overall_gains, 19	counts, 142
counter_overall_loss, 20	duplication, 142
counter_prop_genes_changing, 20	lb, 142
counter_subfun, 20	PhyloRuleDynData, 141
Phylo rules, 21	pos, 142
rule_dyn_limit_changes, 21	ub, 142
phylo.hpp	PhyloRules
get_last_name, 205	-
PHYLO CHECK MISSING, 202	phylo hpp 204
PHYLO_COUNTER_LAMBDA, 202	phylo.hpp, 204
	PhyloRulesDyn
	PhyloRulesDyn phylo.hpp, 204
PHYLO_RULE_DYN_LAMBDA, 202	PhyloRulesDyn phylo.hpp, 204 PhyloStatsCounter
PHYLO_RULE_DYN_LAMBDA, 202 PhyloArray, 203	PhyloRulesDyn phylo.hpp, 204 PhyloStatsCounter phylo.hpp, 204
PHYLO_RULE_DYN_LAMBDA, 202 PhyloArray, 203 PhyloCounter, 203	PhyloRulesDyn phylo.hpp, 204 PhyloStatsCounter phylo.hpp, 204 PhyloSupport
PHYLO_RULE_DYN_LAMBDA, 202 PhyloArray, 203	PhyloRulesDyn phylo.hpp, 204 PhyloStatsCounter phylo.hpp, 204

POS	probability
barraydense-meet.hpp, 181	Node, 136
barraydensecell-meat.hpp, 183	push_back
pos	PhyloCounterData, 140
PhyloRuleDynData, 142	
PowerSet	README.md, 242
PowerSet < Array_Type, Data_Rule_Type >, 144	reduced_sequence
PowerSet < Array_Type, Data_Rule_Type >, 143	Geese, 113
\sim PowerSet, 145	rengine
add_rule, 145	Flock, 98
begin, 145	reserve
calc, 146	BArray< Cell_Type, Data_Type >, 37
coordinates_free, 147	BArrayDense < Cell_Type, Data_Type >, 57
coordinates_locked, 147	FreqTable < T >, 101
data, 148	PhyloCounterData, 140
EmptyArray, 148	reset
end, 146	PowerSet < Array_Type, Data_Rule_Type >, 147
get_data, 146	reset_array
get_data_ptr, 146	StatsCounter< Array_Type, Data_Type >, 157
init_support, 146	Support< Array_Type, Data_Counter_Type
M, 148	Data_Rule_Type, Data_Rule_Dyn_Type >
N, 148	165, 166
operator[], 147	resize
PowerSet, 144	BArray< Cell_Type, Data_Type >, 37
reset, 147	BArrayDense< Cell_Type, Data_Type >, 58
rules, 148	Entries < Cell_Type >, 91
rules_deleted, 149	return
size, 147	support-meat.hpp, 237
predict	rm_cell
Geese, 109	BArray < Cell_Type, Data_Type >, 38
predict_backend	BArrayDense < Cell_Type, Data_Type >, 58
Geese, 110	support-meat.hpp, 231
predict_exhaust	ROW
Geese, 110	barray-meat-operators.hpp, 174
predict_exhaust_backend	barray-meat.hpp, 176
Geese, 110	barraydense-meet.hpp, 181
predict_sim	row
Geese, 110	BArray Cell_Type, Data_Type >, 38
print	BArrayDense < Cell_Type, Data_Type >, 58
BArray< Cell_Type, Data_Type >, 37	Row_type
BArrayDense < Cell_Type, Data_Type >, 57	typedefs.hpp, 240
Flock, 97	Rule
FreqTable $<$ T $>$, 100	Rule < Array_Type, Data_Type >, 150
Geese, 110	Rule < Array_Type, Data_Type >, 149
Model< Array_Type, Data_Counter_Type,	~Rule, 150
Data_Rule_Type, Data_Rule_Dyn_Type >,	D, 151
124	operator(), 151
Support< Array_Type, Data_Counter_Type,	Rule, 150
Data_Rule_Type, Data_Rule_Dyn_Type >,	rule_dyn_limit_changes
165	Phylo rules, 21
print_observed_counts	rule_fun_default rules-bones.hpp, 222
Geese, 111	• •
print_stats	Rule_fun_type typedefs.hpp, 241
Model< Array_Type, Data_Counter_Type,	RULE_FUNCTION
Data_Rule_Type, Data_Rule_Dyn_Type >,	barry.hpp, 189
124	geese-bones.hpp, 212
printf_barry	RULE_LAMBDA
barry-configuration.hpp, 186	barry.hpp, 189
	υαιτιχιτιρρ, του

Rules	Model < Array_Type, Data_Counter_Type,
Rules < Array_Type, Data_Type >, 152	Data_Rule_Type, Data_Rule_Dyn_Type >,
rules	125
PowerSet < Array_Type, Data_Rule_Type >, 148	Support< Array_Type, Data_Counter_Type,
support-meat.hpp, 237	Data_Rule_Type, Data_Rule_Dyn_Type >,
Rules < Array_Type, Data_Type >, 151	166
\sim Rules, 152 add_rule, 153	set_seed Flock, 97
clear, 153	Geese, 111
get_seq, 153	Model < Array_Type, Data_Counter_Type,
operator(), 154	Data_Rule_Type, Data_Rule_Dyn_Type >,
operator=, 154	126
Rules, 152	shrink_to_fit
size, 154	PhyloCounterData, 140
rules-bones.hpp	simulate
rule_fun_default, 222	Geese, 111
rules_	size
support-meat.hpp, 237	BArrayVector $<$ Cell_Type, Data_Type $>$, 70
rules_deleted	BArrayVector_const< Cell_Type, Data_Type >, 74
PowerSet< Array_Type, Data_Rule_Type >, 149	Counters < Array_Type, Data_Type >, 90
rules_dyn	FreqTable < T >, 101
support-meat.hpp, 237	Model < Array_Type, Data_Counter_Type,
rules_zerodiag	Data_Rule_Type, Data_Rule_Dyn_Type >,
network.hpp, 199	126 PhyloCounterData, 140
sample	PowerSet < Array_Type, Data_Rule_Type >, 147
Model< Array_Type, Data_Counter_Type,	Rules < Array_Type, Data_Type >, 154
Data_Rule_Type, Data_Rule_Dyn_Type >,	size_unique
124	Model< Array_Type, Data_Counter_Type,
sequence	Data_Rule_Type, Data_Rule_Dyn_Type >,
Geese, 113	126
set_counters	source
Model< Array_Type, Data_Counter_Type,	Entries < Cell_Type >, 92
Data_Rule_Type, Data_Rule_Dyn_Type >,	states
125	NodeData, 138
StatsCounter < Array_Type, Data_Type >, 158	Statistical Models, 9
Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,	stats_bank
166	support-meat.hpp, 237
set_data	StatsCounter StatsCounter < Array_Type, Data_Type >, 156
BArray< Cell_Type, Data_Type >, 38	StatsCounter< Array_Type, Data_Type >, 155
BArrayDense< Cell Type, Data Type >, 58	~StatsCounter, 156
set_keygen	add_counter, 156
Model < Array_Type, Data_Counter_Type,	count all, 157
Data_Rule_Type, Data_Rule_Dyn_Type >,	count_current, 157
125	count_init, 157
set_rengine	get_counters, 157
Model< Array_Type, Data_Counter_Type,	reset_array, 157
Data_Rule_Type, Data_Rule_Dyn_Type >,	set_counters, 158
125	StatsCounter, 156
set_rules Model	store_psets
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,	Model < Array_Type, Data_Counter_Type,
125	Data_Rule_Type, Data_Rule_Dyn_Type >,
Support< Array_Type, Data_Counter_Type,	126 subtree_prob
Data_Rule_Type, Data_Rule_Dyn_Type >,	Node, 136
166	Support
set_rules_dyn	Support< Array_Type, Data_Counter_Type,

Data_Rule_Type, Data_Rule_Dyn_Type >, 161	support_size Flock, 97
Support < Array_Type, Data_Counter_Type, Data_Rule_1	Type, Geese, 111
Data_Rule_Dyn_Type >, 158	Model< Array_Type, Data_Counter_Type
~Support, 162	Data_Rule_Type, Data_Rule_Dyn_Type >
add_counter, 162	126
add_rule, 162	SUPPORT TEMPLATE
add_rule_dyn, 163	support-meat.hpp, 230, 232–235
calc, 163	SUPPORT TEMPLATE ARGS
change_stats, 166	support-meat.hpp, 230
coordinates_free, 167	SUPPORT TYPE
coordinates_locked, 167	support-meat.hpp, 230
current_stats, 167	swap_cells
delete_counters, 167	BArray< Cell_Type, Data_Type >, 38
delete_rules, 167	BArrayDense < Cell_Type, Data_Type >, 59
delete_rules_dyn, 168	swap_cols
eval rules dyn, 163	BArray< Cell_Type, Data_Type >, 39
get counters, 164	BArrayDense < Cell_Type, Data_Type >, 59
get counts, 164	swap_rows
get_counts_ptr, 164	BArray < Cell_Type, Data_Type >, 39
	BArrayDense < Cell_Type, Data_Type >, 59
get_current_stats, 164 get_data, 164	DAITayDelise Cell_Type, Data_Type >, 59
- -	target
get_rules, 165	Entries< Cell_Type >, 92
get_rules_dyn, 165	toggle_cell
init_support, 165	
M, 168	BArray Cell_Type, Data_Type >, 39
max_num_elements, 168	BArrayDense < Cell_Type, Data_Type >, 59
N, 168	toggle_lock
print, 165	BArray Cell_Type, Data_Type >, 39
reset_array, 165, 166	BArrayDense < Cell_Type, Data_Type >, 60
set_counters, 166	transpose
set_rules, 166	BArray< Cell_Type, Data_Type >, 39
set_rules_dyn, 166	BArrayDense < Cell_Type, Data_Type >, 60
Support, 161	TWO
support-meat.hpp	CHECK, 24
array_bank, 235	EXISTS, 26
BARRY_SUPPORT_MEAT_HPP, 229	typedefs.hpp
calc_backend, 230	Col_type, 240
cfree, 235	Counter_fun_type, 240
counters, 235	Counts_type, 240
counters_, 235	MapVec_type, 240
delete_counters, 236	Row_type, 240
delete_rules, 236	Rule_fun_type, 241
delete_rules_dyn, 236	uint, 241
else, 236	vec_equal, 241
f_, 236	vec_equal_approx, 241
for, 231	vec_inner_prod, 242
if, 231	
insert_cell, 231	ub
return, 237	PhyloRuleDynData, 142
rm_cell, 231	uint
rules, 237	typedefs.hpp, 241
rules_, 237	UKNOWN
rules_dyn, 237	EXISTS, 26
stats bank, 237	update_annotations
-	Geese, 111
SUPPORT_TEMPLATE, 230, 232–235	update_normalizing_constant
SUPPORT_TEMPLATE_ARGS, 230	model-meat.hpp, 209
SUPPORT_TYPE, 230	

```
val
     Entries < Cell_Type >, 92
value
     \mathsf{Cell} \! < \mathsf{Cell} \! \_ \mathsf{Type} >, \textcolor{red}{\textbf{78}}
vec_diff
     geese-bones.hpp, 212
vec_equal
     typedefs.hpp, 241
vec_equal_approx
     typedefs.hpp, 241
vec_inner_prod
     typedefs.hpp, 242
vec Hasher < T >, 169
     operator(), 169
vector_caster
     geese-bones.hpp, 212
vertex_attr
     NetworkData, 130
visited
     {\sf BArray}{<}\ {\sf Cell\_Type},\ {\sf Data\_Type}>,\ {\sf 40}
     BArrayDense < Cell_Type, Data_Type >, 61
     Cell< Cell_Type >, 78
     Node, 136
ZERO_CELL
     barraydense-meet.hpp, 181
zero_col
     BArray< Cell_Type, Data_Type >, 40
     BArrayDense < Cell_Type, Data_Type >, 60
zero_row
     BArray< Cell_Type, Data_Type >, 40
     BArrayDense < Cell_Type, Data_Type >, 60
```