barry: Your go-to motif accountant 0.0-1

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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
5.1.1 Detailed Description	9
5.2 Statistical Models	9
5.2.1 Detailed Description	10
5.3 Network counters	10
5.3.1 Detailed Description	10
5.3.2 Function Documentation	11
5.3.2.1 counter_absdiff()	11
5.3.2.2 counter_ctriads()	11
5.3.2.3 counter_degree()	11
5.3.2.4 counter_density()	11
5.3.2.5 counter_diff()	12
5.3.2.6 counter_edges()	12
5.3.2.7 counter_idegree()	12
5.3.2.8 counter_idegree15()	12
5.3.2.9 counter isolates()	12
5.3.2.10 counter_istar2()	13
5.3.2.11 counter_mutual()	13
5.3.2.12 counter_nodecov()	13
5.3.2.13 counter_nodeicov()	13
5.3.2.14 counter_nodematch()	13
5.3.2.15 counter_nodeocov()	14
5.3.2.16 counter_odegree()	14
5.3.2.17 counter_odegree15()	14
5.3.2.18 counter_ostar2()	14
5.3.2.19 counter_ttriads()	14
5.3.2.20 NETWORK_COUNTER()	15
5.4 Phylo counters	15
5.4.1 Detailed Description	15
5.4.2 Function Documentation	16
5.4.2 Function Documentation	16
5.4.2.2 counter_cogain()	16

5.4.2.3 counter_gains()	1
5.4.2.4 counter_gains_k_offspring()	1
5.4.2.5 counter_genes_changing()	1
5.4.2.6 counter_longest()	1
5.4.2.7 counter_loss()	1
5.4.2.8 counter_maxfuns()	1
5.4.2.9 counter_neofun()	1
5.4.2.10 counter_neofun_a2b()	1
5.4.2.11 counter_overall_changes()	19
5.4.2.12 counter_overall_gains()	19
5.4.2.13 counter_overall_loss()	19
5.4.2.14 counter_prop_genes_changing()	19
5.4.2.15 counter_subfun()	2
5.5 Phylo rules	2
5.5.1 Detailed Description	2
5.5.2 Function Documentation	2
5.5.2.1 rule_dyn_limit_changes()	2
	_
6 Namespace Documentation	2
6.1 barry Namespace Reference	
6.1.1 Detailed Description	
6.2 barry::counters Namespace Reference	
6.2.1 Detailed Description	
6.3 barry::counters::network Namespace Reference	
6.4 barry::counters::phylo Namespace Reference	
6.5 CHECK Namespace Reference	
6.5.1 Detailed Description	2
6.5.2 Variable Documentation	
6.5.2.1 BOTH	
6.5.2.2 NONE	2
6.5.2.3 ONE	
6.5.2.4 TWO	2
6.6 EXISTS Namespace Reference	2
6.6.1 Detailed Description	2
6.6.2 Variable Documentation	2
6.6.2.1 AS_ONE	2
6.6.2.2 AS_ZERO	2
6.6.2.3 BOTH	2
6.6.2.4 NONE	2
6.6.2.5 ONE	2
6.6.2.6 TWO	2
6.6.2.7 UKNOWN	2

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
<b>7.1.2.2 BArray()</b> [2/6]	30
<b>7.1.2.3 BArray()</b> [3/6]	30
7.1.2.4 BArray() [4/6]	31
<b>7.1.2.5 BArray()</b> [5/6]	31
<b>7.1.2.6 BArray()</b> [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
<b>7.4.2.3 BArrayDense()</b> [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]	56
7.4.3.25 operator-=() [1/3]	56

7.4.3.26 operator-=() [2/3]	56
<b>7.4.3.27 operator-=()</b> [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 \; BArray Dense Cell < 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+=()	
7.5.3.4 operator-=()	
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	
7.6.1 Detailed Description	
7.6.2 Constructor & Destructor Documentation	65

7.6.2.1 BArrayDenseCell_const()	65
7.6.2.2 ~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 BArrayVector_const< Cell_Type, Data_Type > Class Template Reference	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 ~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
<b>7.9.2.2 Cell()</b> [2/7]	75
7.9.2.3 ~Cell()	75
<b>7.9.2.4 Cell()</b> [3/7]	76
7.9.2.5 Cell() [4/7]	76
<b>7.9.2.6 Cell()</b> [5/7]	76
<b>7.9.2.7 Cell()</b> [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
<b>7.9.3.3 add()</b> [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() [1.74]	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	
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	
7.12.2 Constructor & Destructor Documentation	
7.12.2.1 Counters() [1/3]	 87
7.12.2.2 ~Counters()	 . 87
<b>7.12.2.3 Counters()</b> [2/3]	
<b>7.12.2.4 Counters()</b> [3/3]	
7.12.3 Member Function Documentation	
<b>7.12.3.1 add_counter()</b> [1/3]	
7.12.3.2 add_counter() [2/3]	 . 88
<b>7.12.3.3 add_counter()</b> [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[]()	
7.12.3.8 size()	
7.13 Entries < Cell_Type > Class Template Reference	
7.13.1 Detailed Description	
7.13.2 Constructor & Destructor Documentation	
7.13.2.1 Entries() [1/2]	
7.13.2.2 Entries() [2/2]	
7.13.2.3 ~Entries()	
7.13.3 Member Function Documentation	
7.13.3.1 resize()	
7.13.4 Member Data Documentation	
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()	
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()	110
7.16.3.30 print()	111

7.16.3.31 print_observed_counts()	1	111
7.16.3.32 set_seed()	1	111
7.16.3.33 simulate()	1	111
7.16.3.34 support_size()	1	111
7.16.3.35 update_annotations()	1	112
7.16.4 Member Data Documentation	1	112
7.16.4.1 delete_rengine	1	112
7.16.4.2 delete_support	1	112
7.16.4.3 initialized	1	112
7.16.4.4 map_to_nodes	1	112
7.16.4.5 nfunctions	1	113
7.16.4.6 nodes	1	113
7.16.4.7 reduced_sequence	1	113
7.16.4.8 sequence	1	113
7.17 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class Tell		
plate Reference		
7.17.1 Detailed Description		
7.17.2 Constructor & Destructor Documentation		
7.17.2.1 Model() [1/3]		
<b>7.17.2.2 Model()</b> [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]		
<b>7.17.3.3</b> add_counter() [2/3]		
<b>7.17.3.4</b> add_counter() [3/3]		
7.17.3.5 add_rule() [1/3]		
<b>7.17.3.6 add_rule()</b> [2/3]		
<b>7.17.3.7 add_rule()</b> [3/3]		
7.17.3.8 add_rule_dyn() [1/3]		
7.17.3.9 add_rule_dyn() [2/3]		
<b>7.17.3.10 add_rule_dyn()</b> [3/3]		
7.17.3.11 colnames()		
7.17.3.12 gen_key()		
7.17.3.13 get_counters()		
7.17.3.14 get_norm_const()		
7.17.3.15 get_pset()		
7.17.3.16 get_pset_stats()		
7.17.3.17 get_rengine()		
7.17.3.18 get_rules()		
7.17.3.19 get_rules_dyn()	1	122

7.17.3.20 get_support()	 122
<b>7.17.3.21 likelihood()</b> [1/3]	 122
<b>7.17.3.22 likelihood()</b> [2/3]	 122
<b>7.17.3.23 likelihood()</b> [3/3]	 122
7.17.3.24 likelihood_total()	 123
7.17.3.25 nterms()	 123
7.17.3.26 operator=()	 123
7.17.3.27 print()	 123
7.17.3.28 print_stats()	 123
7.17.3.29 sample() [1/2]	 124
7.17.3.30 sample() [2/2]	 124
7.17.3.31 set_counters()	 124
7.17.3.32 set_keygen()	 124
7.17.3.33 set_rengine()	 124
7.17.3.34 set_rules()	 125
7.17.3.35 set_rules_dyn()	 125
7.17.3.36 set_seed()	 125
7.17.3.37 size()	 125
7.17.3.38 size_unique()	 125
7.17.3.39 store_psets()	 126
7.17.3.40 support_size()	 126
7.18 NetCounterData Class Reference	 126
7.18.1 Detailed Description	 126
7.18.2 Constructor & Destructor Documentation	 126
7.18.2.1 NetCounterData() [1/2]	 127
7.18.2.2 NetCounterData() [2/2]	 127
7.18.2.3 ∼NetCounterData()	 127
7.18.3 Member Data Documentation	 127
7.18.3.1 indices	 127
7.18.3.2 numbers	 127
7.19 NetworkData Class Reference	 128
7.19.1 Detailed Description	 128
7.19.2 Constructor & Destructor Documentation	 128
7.19.2.1 NetworkData() [1/3]	 128
7.19.2.2 NetworkData() [2/3]	 128
7.19.2.3 NetworkData() [3/3]	 129
7.19.2.4 ~NetworkData()	 129
7.19.3 Member Data Documentation	 129
7.19.3.1 directed	 129
7.19.3.2 vertex_attr	 130
7.20 Node Class Reference	 130
7.20.1 Detailed Description	 131

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

7.22.3.6 operator()()	19
7.22.3.7 push_back()	9
7.22.3.8 reserve()	19
7.22.3.9 shrink_to_fit()	19
7.22.3.10 size()	0
7.23 PhyloRuleDynData Class Reference	0
7.23.1 Detailed Description	0
7.23.2 Constructor & Destructor Documentation	0
7.23.2.1 PhyloRuleDynData()	0
7.23.2.2 ∼PhyloRuleDynData()	1
7.23.3 Member Data Documentation	⊦1
7.23.3.1 counts	1
7.23.3.2 duplication	
7.23.3.3 lb	
7.23.3.4 pos	∤1
7.23.3.5 ub	⊦1
7.24 PowerSet< Array_Type, Data_Rule_Type > Class Template Reference	
7.24.1 Detailed Description	3
7.24.2 Constructor & Destructor Documentation	
7.24.2.1 PowerSet() [1/3]	3
7.24.2.2 PowerSet() [2/3]	3
7.24.2.3 PowerSet() [3/3]	
7.24.2.4 ~PowerSet()	
7.24.3 Member Function Documentation	
7.24.3.1 add_rule() [1/3]	
7.24.3.2 add_rule() [2/3]	
7.24.3.3 add_rule() [3/3]14	4
7.24.3.4 begin()	
7.24.3.5 calc()	
7.24.3.6 end()	
7.24.3.7 get_data()	
7.24.3.8 get_data_ptr()	
7.24.3.9 init_support()	
7.24.3.10 operator[]()	
7.24.3.11 reset()	
7.24.3.12 size()	
7.24.4 Member Data Documentation	
7.24.4.1 coordinates_free	
7.24.4.2 coordinates_locked	
7.24.4.3 data	
7.24.4.4 EmptyArray	
7.24.4.5 M	-/

7.24.4.6 N	147
7.24.4.7 rules	148
7.24.4.8 rules_deleted	148
7.25 Rule < Array_Type, Data_Type > Class Template Reference	148
7.25.1 Detailed Description	149
7.25.2 Constructor & Destructor Documentation	149
7.25.2.1 Rule() [1/2]	149
<b>7.25.2.2 Rule()</b> [2/2]	149
7.25.2.3 ∼Rule()	149
7.25.3 Member Function Documentation	150
7.25.3.1 D()	150
7.25.3.2 operator()()	150
7.26 Rules< Array_Type, Data_Type > Class Template Reference	150
7.26.1 Detailed Description	151
7.26.2 Constructor & Destructor Documentation	151
7.26.2.1 Rules() [1/2]	151
7.26.2.2 Rules() [2/2]	151
7.26.2.3 ∼Rules()	152
7.26.3 Member Function Documentation	152
7.26.3.1 add_rule() [1/3]	152
<b>7.26.3.2 add_rule()</b> [2/3]	152
<b>7.26.3.3 add_rule()</b> [3/3]	152
7.26.3.4 clear()	152
7.26.3.5 get_seq()	152
7.26.3.6 operator()()	153
7.26.3.7 operator=()	153
7.26.3.8 size()	154
7.27 StatsCounter< Array_Type, Data_Type > Class Template Reference	154
7.27.1 Detailed Description	154
7.27.2 Constructor & Destructor Documentation	155
7.27.2.1 StatsCounter() [1/2]	155
<b>7.27.2.2 StatsCounter()</b> [2/2]	155
7.27.2.3 ∼StatsCounter()	155
7.27.3 Member Function Documentation	155
7.27.3.1 add_counter() [1/2]	155
7.27.3.2 add_counter() [2/2]	156
7.27.3.3 count_all()	156
7.27.3.4 count_current()	156
7.27.3.5 count_init()	156
7.27.3.6 get_counters()	156
7.27.3.7 reset_array()	156
7.27.3.8 set_counters()	157

7.28 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class Tem	
plate Reference	
7.28.1 Detailed Description	
7.28.2 Constructor & Destructor Documentation	
7.28.2.1 Support() [1/3]	
7.28.2.2 Support() [2/3]	
7.28.2.3 Support() [3/3]	
7.28.2.4 ~Support()	
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]	
<b>7.28.3.6 add_rule_dyn()</b> [2/2]	
7.28.3.7 calc()	
7.28.3.8 eval_rules_dyn()	
7.28.3.9 get_counters()	
7.28.3.10 get_counts()	
7.28.3.11 get_counts_ptr()	
7.28.3.12 get_current_stats()	
7.28.3.13 get_data()	
7.28.3.14 get_rules()	
7.28.3.15 get_rules_dyn()	
7.28.3.16 init_support()	
7.28.3.17 print()	. 164
7.28.3.18 reset_array() [1/2]	
<b>7.28.3.19 reset_array()</b> [2/2]	. 165
7.28.3.20 set_counters()	. 165
7.28.3.21 set_rules()	. 165
7.28.3.22 set_rules_dyn()	. 165
7.28.4 Member Data Documentation	. 165
7.28.4.1 change_stats	. 166
7.28.4.2 coordinates_free	. 166
7.28.4.3 coordinates_locked	. 166
7.28.4.4 current_stats	. 166
7.28.4.5 delete_counters	. 166
7.28.4.6 delete_rules	. 167
7.28.4.7 delete_rules_dyn	. 167
7.28.4.8 M	. 167
7.28.4.9 max_num_elements	. 167
7.28.4.10 N	. 167

	7.29 vecHasher $<$ T $>$ Struct Template Reference	168
	7.29.1 Detailed Description	168
	7.29.2 Member Function Documentation	168
	7.29.2.1 operator()()	168
8 I	File Documentation	169
	8.1 include/barry/barray-bones.hpp File Reference	169
	8.1.1 Macro Definition Documentation	170
	8.1.1.1 BARRAY_BONES_HPP	170
	8.2 include/barry/barray-iterator.hpp File Reference	170
	8.3 include/barry/barray-meat-operators.hpp File Reference	171
	8.3.1 Macro Definition Documentation	172
	8.3.1.1 BARRY_BARRAY_MEAT_OPERATORS_HPP	172
	8.3.1.2 COL	172
	8.3.1.3 ROW	172
	8.3.2 Function Documentation	172
	8.3.2.1 checkdim_()	172
	8.4 include/barry/barray-meat.hpp File Reference	173
	8.4.1 Macro Definition Documentation	174
	8.4.1.1 COL	174
	8.4.1.2 ROW	174
	8.5 include/barry/barraycell-bones.hpp File Reference	174
	8.6 include/barry/barraycell-meat.hpp File Reference	175
	8.7 include/barry/barraydense-bones.hpp File Reference	176
	8.8 include/barry/barraydense-meet.hpp File Reference	177
	8.8.1 Macro Definition Documentation	178
	8.8.1.1 BARRY_BARRAYDENSE_MEAT_HPP	178
	8.8.1.2 COL	179
	8.8.1.3 POS	179
	8.8.1.4 ROW	179
	8.8.1.5 ZERO_CELL	179
	8.9 include/barry/barraydensecell-bones.hpp File Reference	179
	8.10 include/barry/barraydensecell-meat.hpp File Reference	180
	8.10.1 Macro Definition Documentation	181
	8.10.1.1 BARRY_BARRAYDENSECELL_MEAT_HPP	181
	8.10.1.2 POS	181
	8.11 include/barry/barrayvector-bones.hpp File Reference	181
	8.12 include/barry/barrayvector-meat.hpp File Reference	182
	8.12.1 Macro Definition Documentation	183
	8.12.1.1 BARRY_BARRAYVECTOR_MEAT_HPP	183
	8.13 include/barry/barry-configuration.hpp File Reference	183
	8.13.1 Macro Definition Documentation	184

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
8.21.1 Macro Definition Documentation
8 21 1 1 PHYLO CHECK MISSING

8.21.1.2 PHYLO_COUNTER_LAMBDA
8.21.1.3 PHYLO_RULE_DYN_LAMBDA
8.21.2 Typedef Documentation
8.21.2.1 PhyloArray
8.21.2.2 PhyloCounter
8.21.2.3 PhyloCounters
8.21.2.4 PhyloModel
8.21.2.5 PhyloPowerSet
8.21.2.6 PhyloRule
8.21.2.7 PhyloRuleData
8.21.2.8 PhyloRuleDyn
8.21.2.9 PhyloRules
8.21.2.10 PhyloRulesDyn
8.21.2.11 PhyloStatsCounter
8.21.2.12 PhyloSupport
8.21.3 Function Documentation
8.21.3.1 get_last_name()
8.22 include/barry/model-bones.hpp File Reference
8.22.1 Function Documentation
8.22.1.1 keygen_default()
8.23 include/barry/model-meat.hpp File Reference
8.23.1 Macro Definition Documentation
8.23.1.1 MODEL_TEMPLATE
8.23.1.2 MODEL_TEMPLATE_ARGS
8.23.1.3 MODEL_TYPE
8.23.2 Function Documentation
8.23.2.1 likelihood_()
8.23.2.2 MODEL_TEMPLATE() [1/2]
8.23.2.3 MODEL_TEMPLATE() [2/2]
8.23.2.4 update_normalizing_constant()
8.24 include/barry/models/geese.hpp File Reference
8.25 include/barry/models/geese/flock-bones.hpp File Reference
8.26 include/barry/models/geese/flock-meet.hpp File Reference
8.27 include/barry/models/geese/geese-bones.hpp File Reference
8.27.1 Macro Definition Documentation
8.27.1.1 INITIALIZED
8.27.2 Function Documentation
8.27.2.1 keygen_full()
8.27.2.2 RULE_FUNCTION()
8.27.2.3 vec_diff()
8.27.2.4 vector_caster()
8.28 include/barry/models/geese/geese-meat-constructors hop File Reference 210

$8.29\ include/barry/models/geese/geese-meat-likelihood. hpp\ File\ Reference\ .\ .\ .\ .\ .\ .\ .\ .\ .\ .\ .\ .\ .\$
$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
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
8.34 include/barry/models/geese/geese-meat-simulate.hpp File Reference
8.35 include/barry/models/geese/geese-meat.hpp File Reference
8.36 include/barry/models/geese/geese-node-bones.hpp File Reference
8.37 include/barry/powerset-bones.hpp File Reference
8.38 include/barry/powerset-meat.hpp File Reference
8.39 include/barry/rules-bones.hpp File Reference
8.39.1 Function Documentation
8.39.1.1 rule_fun_default()
8.40 include/barry/rules-meat.hpp File Reference
8.41 include/barry/statscounter-bones.hpp File Reference
8.42 include/barry/statscounter-meat.hpp File Reference
8.43 include/barry/statsdb.hpp File Reference
8.44 include/barry/support-bones.hpp File Reference
8.45 include/barry/support-meat.hpp File Reference
8.45.1 Macro Definition Documentation
8.45.1.1 BARRY_SUPPORT_MEAT_HPP
8.45.1.2 SUPPORT_TEMPLATE
8.45.1.3 SUPPORT_TEMPLATE_ARGS
8.45.1.4 SUPPORT_TYPE
8.45.2 Function Documentation
8.45.2.1 calc_backend()
8.45.2.2 for()
8.45.2.3 if() [1/3]
8.45.2.4 if() [2/3]
8.45.2.5 if() [3/3]
8.45.2.6 insert_cell()
8.45.2.7 rm_cell()
8.45.2.8 SUPPORT_TEMPLATE() [1/17]
8.45.2.9 SUPPORT_TEMPLATE() [2/17]
8.45.2.10 SUPPORT_TEMPLATE() [3/17]
8.45.2.11 SUPPORT_TEMPLATE() [4/17]
8.45.2.12 SUPPORT_TEMPLATE() [5/17]
8.45.2.13 SUPPORT_TEMPLATE() [6/17]
8.45.2.14 SUPPORT_TEMPLATE() [7/17]
8.45.2.15 SUPPORT_TEMPLATE() [8/17]
<b>8.45.2.16 SUPPORT_TEMPLATE()</b> [9/17]
8.45.2.17 SUPPORT_TEMPLATE() [10/17]

8.45.2.18 SUPPORT_TEMPLATE() [11/17]	231
<b>8.45.2.19 SUPPORT_TEMPLATE()</b> [12/17]	231
<b>8.45.2.20 SUPPORT_TEMPLATE()</b> [13/17]	231
8.45.2.21 SUPPORT_TEMPLATE() [14/17]	231
<b>8.45.2.22 SUPPORT_TEMPLATE()</b> [15/17]	231
<b>8.45.2.23 SUPPORT_TEMPLATE()</b> [16/17]	232
8.45.2.24 SUPPORT_TEMPLATE() [17/17]	232
8.45.3 Variable Documentation	232
8.45.3.1 array_bank	232
8.45.3.2 cfree	232
8.45.3.3 counters	232
8.45.3.4 counters	233
8.45.3.5 delete_counters	233
8.45.3.6 delete_rules	233
8.45.3.7 delete_rules_dyn	233
8.45.3.8 else	233
8.45.3.9 f	234
8.45.3.10 return	234
8.45.3.11 rules	234
8.45.3.12 rules	234
8.45.3.13 rules_dyn	234
8.45.3.14 stats_bank	235
8.46 include/barry/typedefs.hpp File Reference	235
8.46.1 Typedef Documentation	237
8.46.1.1 Col_type	237
8.46.1.2 Counter_fun_type	237
8.46.1.3 Counts_type	237
8.46.1.4 MapVec_type	237
8.46.1.5 Row_type	238
8.46.1.6 Rule_fun_type	238
8.46.1.7 uint	238
8.46.2 Function Documentation	238
8.46.2.1 vec_equal()	238
8.46.2.2 vec_equal_approx()	239
8.46.2.3 vec_inner_prod()	239
8.47 README.md File Reference	239
Index	241

## 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																							9
Statistical Models																							ç
Network counters																							10
Phylo counters																							15
Phylo rules																							20

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
BArrayCell < Cell_Type, Data_Type >
BArrayCell_const< Cell_Type, Data_Type >
BArrayDense < Cell_Type, Data_Type >
Baseline class for binary arrays
BArrayDenseCell< Cell_Type, Data_Type >
BArrayDenseCell_const< Cell_Type, Data_Type >
BArrayVector< Cell_Type, Data_Type >
Row or column of a BArray 6
BArrayVector_const < Cell_Type, Data_Type >
Cell< Cell_Type >
Entries in BArray. For now, it only has two members:
ConstBArrayRowlter < Cell_Type, Data_Type >
Counter< Array_Type, Data_Type >
A counter function based on change statistics
Counters < Array_Type, Data_Type >
Vector of counters
Entries < Cell_Type >
A wrapper class to store source, target, val from a BArray object 9
Flock
A Flock is a group of Geese
FreqTable < T >
Database of statistics
Geese
Annotated Phylo Model
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:
NetCounterData
Data class used to store arbitrary uint or double vectors
NetworkData
Data class for Networks
Node
A single node for the model

6 Class Index

NodeData	
Data definition for the PhyloArray class	136
PhyloCounterData	137
PhyloRuleDynData	140
PowerSet< Array_Type, Data_Rule_Type >	
Powerset of a binary array	142
Rule < Array_Type, Data_Type >	
Rule for determining if a cell should be included in a sequence	148
Rules < Array_Type, Data_Type >	
Vector of objects of class Rule	150
StatsCounter< Array_Type, Data_Type >	
Count stats for a single Array	154
Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >	
Compute the support of sufficient statistics	157
vecHasher< T >	168

# 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	207
include/barry/models/geese/geese-bones.hpp	208
include/barry/models/geese/geese-meat-constructors.hpp	210
include/barry/models/geese/geese-meat-likelihood.hpp	210
include/barry/models/geese/geese-meat-likelihood_exhaust.hpp	211
include/barry/models/geese/geese-meat-predict.hpp	212
include/barry/models/geese/geese-meat-predict_exhaust.hpp	212
include/barry/models/geese/geese-meat-predict_sim.hpp	213
include/barry/models/geese/geese-meat-simulate.hpp	213
include/barry/models/geese/geese-meat.hpp	214
	214

### **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.

#### 5.3.1 Detailed Description

Counters for network models.

5.3 Network counters

#### **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 419 of file network.hpp.

#### 5.3.2.2 counter\_ctriads()

Definition at line 322 of file network.hpp.

#### 5.3.2.3 counter\_degree()

Counts number of vertices with a given out-degree.

Definition at line 690 of file network.hpp.

#### 5.3.2.4 counter\_density()

Definition at line 361 of file network.hpp.

12 Module Documentation

#### 5.3.2.5 counter\_diff()

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

Definition at line 461 of file network.hpp.

#### 5.3.2.6 counter\_edges()

Number of edges.

Definition at line 128 of file network.hpp.

### 5.3.2.7 counter\_idegree()

Counts number of vertices with a given in-degree.

Definition at line 604 of file network.hpp.

#### 5.3.2.8 counter\_idegree15()

Definition at line 377 of file network.hpp.

#### 5.3.2.9 counter\_isolates()

Number of isolated vertices.

Definition at line 142 of file network.hpp.

5.3 Network counters

#### 5.3.2.10 counter\_istar2()

Definition at line 210 of file network.hpp.

#### 5.3.2.11 counter\_mutual()

Number of mutual ties.

Definition at line 172 of file network.hpp.

#### 5.3.2.12 counter\_nodecov()

Definition at line 558 of file network.hpp.

#### 5.3.2.13 counter\_nodeicov()

Definition at line 520 of file network.hpp.

#### 5.3.2.14 counter\_nodematch()

Definition at line 578 of file network.hpp.

14 Module Documentation

# 5.3.2.15 counter\_nodeocov()

Definition at line 539 of file network.hpp.

#### 5.3.2.16 counter\_odegree()

Counts number of vertices with a given out-degree.

Definition at line 646 of file network.hpp.

#### 5.3.2.17 counter\_odegree15()

Definition at line 397 of file network.hpp.

#### 5.3.2.18 counter\_ostar2()

Definition at line 228 of file network.hpp.

#### 5.3.2.19 counter\_ttriads()

Definition at line 247 of file network.hpp.

5.4 Phylo counters 15

#### 5.3.2.20 NETWORK\_COUNTER()

Definition at line 503 of file network.hpp.

# 5.4 Phylo counters

Counters for phylogenetic modeling.

#### **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.

16 Module Documentation

#### **Parameters**

counters

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

#### 5.4.2 Function Documentation

#### 5.4.2.1 counter\_co\_opt()

Function co-opting.

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

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

This algorithm implements the change statistic.

Definition at line 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 Phylo counters

#### 5.4.2.3 counter\_gains()

Functional gains for a specific function (nfun).

Definition at line 179 of file phylo.hpp.

#### 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.

18 Module Documentation

#### 5.4.2.7 counter\_loss()

Total count of losses for an specific function.

Definition at line 612 of file phylo.hpp.

#### 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 Phylo counters

#### 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.

#### 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.

20 Module Documentation

#### 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

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()

5.5 Phylo rules 21

```
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
ub	Upper bound

#### Returns

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

Definition at line 1317 of file phylo.hpp.

22 Module Documentation

# **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
- $\bullet \ \ \text{void} \ \ \underline{\text{get\_col\_vec}} \ \ (\text{std::vector} < \ \ \text{Cell\_Type} > *x, \ \underline{\text{uint i, bool check\_bounds=true}}) \ \ \text{const}$
- 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</li>
- 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</li>
- 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]

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

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	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	e 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 > &params={})
- Array\_Type sample (const uint &i, const std::vector< double > &params)
- 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</t> 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 add\_rule\_dyn (Rule\_fun\_type < Array\_Type, Data\_Rule\_Dyn\_Type > count\_fun\_, Data\_Rule\_Dyn
   \_Type \*data\_=nullptr, bool delete\_data\_=false)
- void set\_rules\_dyn (Rules < Array\_Type, Data\_Rule\_Dyn\_Type > \*rules\_)

#### Likelihood functions.

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

#### **Parameters**

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

- double likelihood (const std::vector< double > &params, const uint &i, bool as\_log=false)
- double likelihood (const std::vector< double > &params, const std::vector< double > &target\_, const uint &i, bool as log=false)
- double likelihood\_total (const std::vector< double > &params, bool as\_log=false)

## Extract elements by index

#### **Parameters**

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

- double get norm const (const std::vector< double > &params, const 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 52 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 152 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 gen key()

# 7.17.3.13 get\_counters()

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

# 7.17.3.14 get\_norm\_const()

# 7.17.3.15 get\_pset()

#### 7.17.3.16 get pset stats()

# 7.17.3.17 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
```

# 7.17.3.18 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 ()
```

# 7.17.3.19 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 ()
```

### 7.17.3.20 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 ( )
```

# 7.17.3.21 likelihood() [1/3]

# 7.17.3.22 likelihood() [2/3]

### 7.17.3.23 likelihood() [3/3]

#### 7.17.3.24 likelihood\_total()

#### 7.17.3.25 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]
```

# 7.17.3.26 operator=()

#### 7.17.3.27 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.

# 7.17.3.28 print\_stats()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\longleftrightarrow Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::print_stats ( uint i) const
```

# 7.17.3.29 sample() [1/2]

#### 7.17.3.30 sample() [2/2]

# 7.17.3.31 set\_counters()

# 7.17.3.32 set\_keygen()

# 7.17.3.33 set rengine()

Definition at line 124 of file model-bones.hpp.

### 7.17.3.34 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_ )
```

# 7.17.3.35 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_ )
```

#### 7.17.3.36 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 134 of file model-bones.hpp.

# 7.17.3.37 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]
```

# 7.17.3.38 size\_unique()

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

#### 7.17.3.39 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]
```

# 7.17.3.40 support\_size()

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

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.

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

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.

# 7.19.3.2 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)

7.20 Node Class Reference 131

#### **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.

# 7.20.2.2 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 Node Class Reference 133

# 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.

# 7.20.4.3 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 Node Class Reference 135

#### 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

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

#### **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 ()

158 Class Documentation

Vector of couter functions.

Rules< Array\_Type, Data\_Rule\_Type > \* get\_rules ()

Vector of static rules (cells to iterate).

• Rules< Array\_Type, Data\_Rule\_Dyn\_Type > \* get\_rules\_dyn ()

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

# Resets the support calculator

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

#### **Parameters**

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

- void reset\_array ()
- void reset\_array (const Array\_Type &Array\_)

#### **Manage counters**

#### **Parameters**

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

- void add\_counter (Counter< Array\_Type, Data\_Counter\_Type > \*f\_)
- void 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
- $std::vector < std::pair < uint, uint >> coordinates_free$
- std::vector< std::pair< uint, uint >> coordinates\_locked
- std::vector< std::vector< double >> change stats

160 Class Documentation

### 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.

#### 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]

162 Class Documentation

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

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

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarrow Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::add_rule_\leftarrow dyn ( Rule< Array_Type, Data_Rule_Dyn_Type > * f_- )
```

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

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\longleftrightarrow 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_\longleftrightarrow 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.

#### 7.28.3.8 eval\_rules\_dyn()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\hookleftarrow Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> bool Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::eval_\hookleftarrow rules_dyn ( const std::vector< double > & counts, const uint & i, const uint & j)
```

#### 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.

164 Class Documentation

#### 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
```

# 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

166 Class 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.

#### 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

168 Class Documentation

# 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

# **Chapter 8**

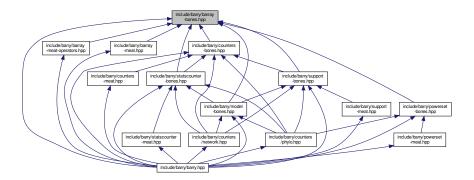
# **File Documentation**

# 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.

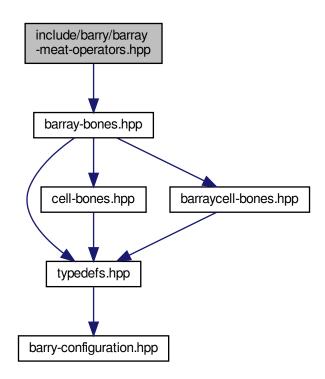
# 8.2 include/barry/barray-iterator.hpp File Reference

### **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.

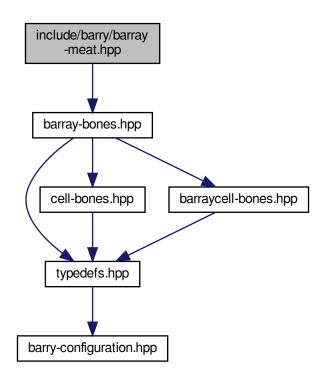
### 8.3.2 Function Documentation

# 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.

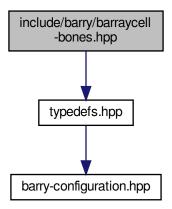
# 8.4.1.2 ROW

```
#define ROW( a \ ) \ \  \mbox{this->el_ij[a]}
```

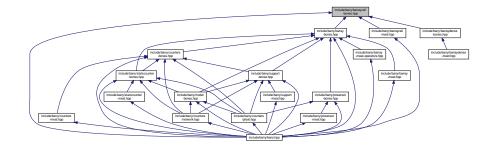
Definition at line 7 of file barray-meat.hpp.

# 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:

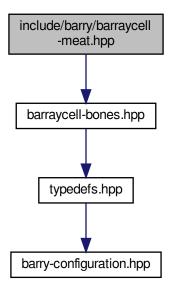


#### **Classes**

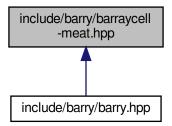
- class BArrayCell
   Cell\_Type, Data\_Type
- class BArrayCell\_const< Cell\_Type, Data\_Type >

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

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



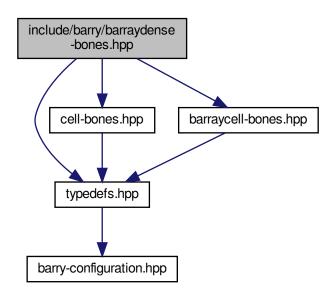
This graph shows which files directly or indirectly include this file:



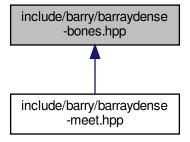
# 8.7 include/barry/barraydense-bones.hpp File Reference

```
#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:



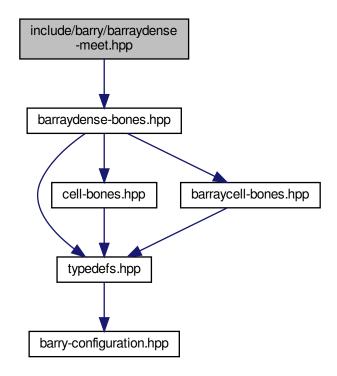
# Classes

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 \ ) \ \  \mbox{this->el_ij[a]}
```

Definition at line 7 of file barraydense-meet.hpp.

# 8.8.1.5 ZERO\_CELL

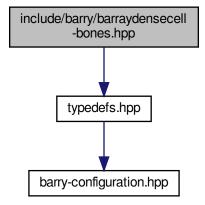
```
#define ZERO_CELL Cell Cell_Type >(static_cast< Cell_Type >(0.0))
```

Definition at line 14 of file barraydense-meet.hpp.

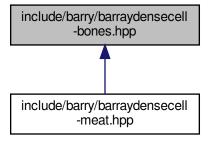
# 8.9 include/barry/barraydensecell-bones.hpp File Reference

```
#include "typedefs.hpp"
```

Include dependency graph for barraydensecell-bones.hpp:



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

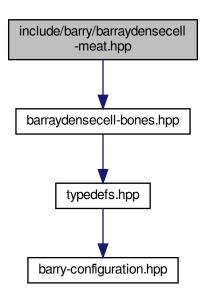


#### **Classes**

- 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.

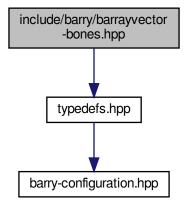
#### 8.10.1.2 POS

```
#define POS(  a \text{,} \\ b \text{ ) (a) + (b) * Array->N }
```

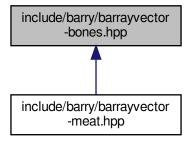
Definition at line 6 of file barraydensecell-meat.hpp.

# 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:

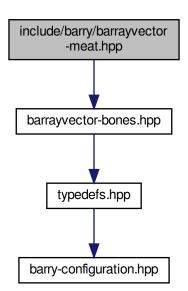


# **Classes**

- 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

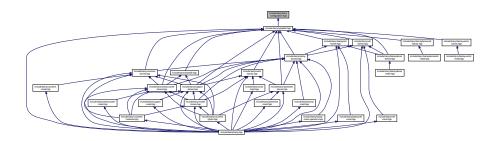
#### 8.12.1.1 BARRY BARRAYVECTOR MEAT HPP

#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

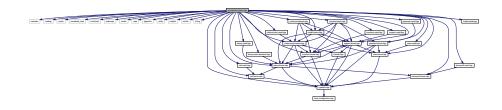
#### 8.13.2.1 Map

```
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

# 8.14.1.5 RULE\_FUNCTION

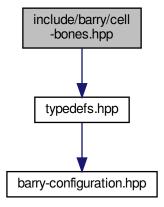
Definition at line 80 of file barry.hpp.

Definition at line 83 of file barry.hpp.

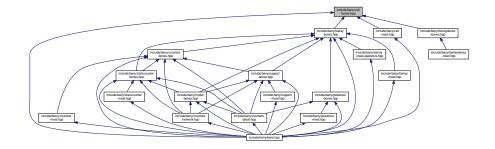
#### 8.14.1.6 RULE\_LAMBDA

# 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:

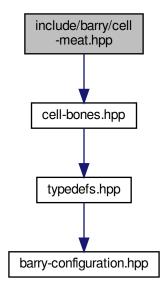


# **Classes**

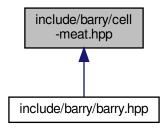
class Cell
 Cell\_Type >
 Entries in BArray. For now, it only has two members:

# 8.16 include/barry/cell-meat.hpp File Reference

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



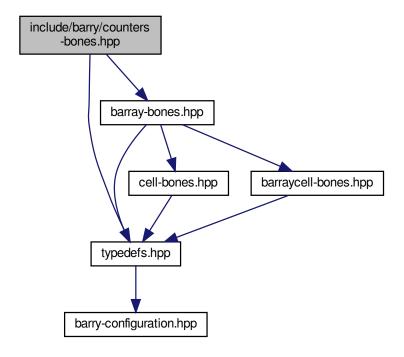
This graph shows which files directly or indirectly include this file:



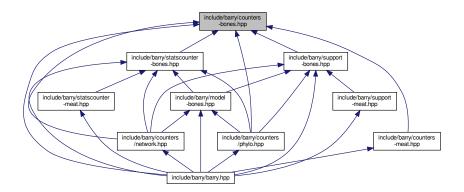
# 8.17 include/barry/col-bones.hpp File Reference

# 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:



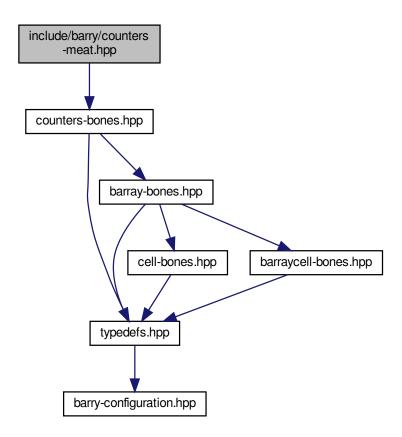
# Classes

- 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:



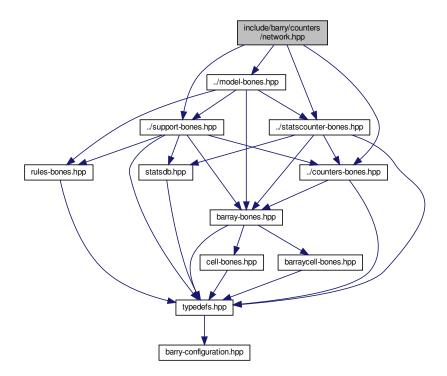
This graph shows which files directly or indirectly include this file:



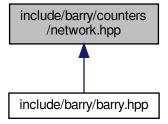
# 8.20 include/barry/counters/network.hpp File Reference

```
#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.

#### 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

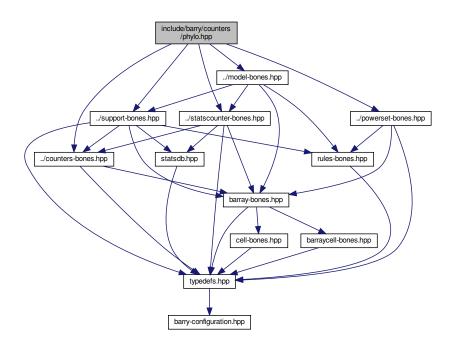
# 8.20.3.1 rules\_zerodiag()

Number of edges.

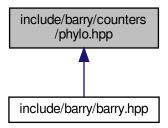
Definition at line 803 of file network.hpp.

# 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

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

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

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

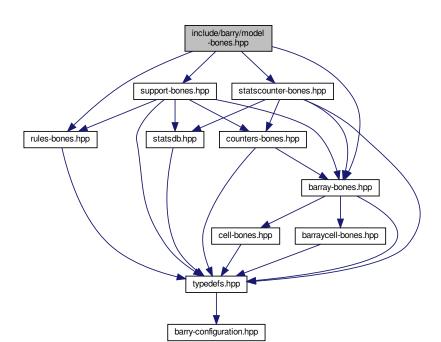
# 8.21.3.1 get\_last\_name()

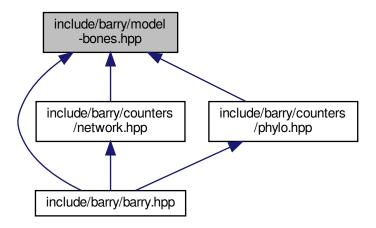
```
std::string get_last_name (
          bool d ) [inline]
```

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:
```





#### **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

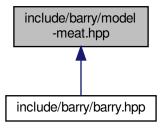
#### 8.22.1.1 keygen\_default()

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 > &params, const Counts\_type &support)
- double likelihood\_ (const std::vector< double > &target\_stats, const std::vector< double > &params, 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

```
#define MODEL_TEMPLATE_ARGS( )
Value:
```

<typename Array\_Type, typename Data\_Counter\_Type,\
typename Data\_Rule\_Type, typename Data\_Rule\_Dyn\_Type>

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

#### 8.23.1.3 MODEL\_TYPE

```
#define MODEL_TYPE( )

Value:
    Model<Array_Type, Data_Counter_Type, Data_Rule_Type, \
    Data_Rule_Dyn_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]

```
MODEL_TEMPLATE (

Model )
```

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:

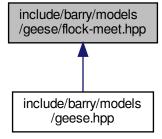


#### **Classes**

· 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)
- bool vec\_diff (const std::vector< unsigned int > &s, const std::vector< unsigned int > &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

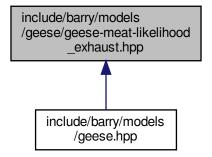
#include "geese-bones.hpp"
Include dependency graph for geese-meat-likelihood.hpp:



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:

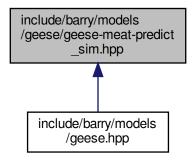


# 8.32 include/barry/models/geese/geese-meat-predict\_exhaust.hpp File Reference



# 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

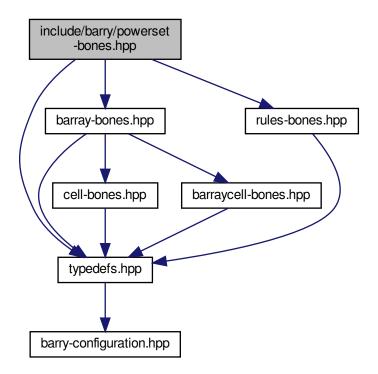
• class Node

A single node for the model.

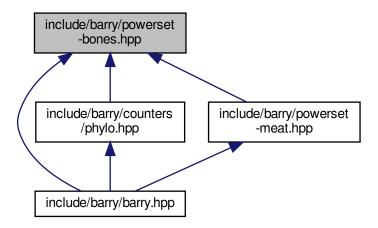
# 8.37 include/barry/powerset-bones.hpp File Reference

```
#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:

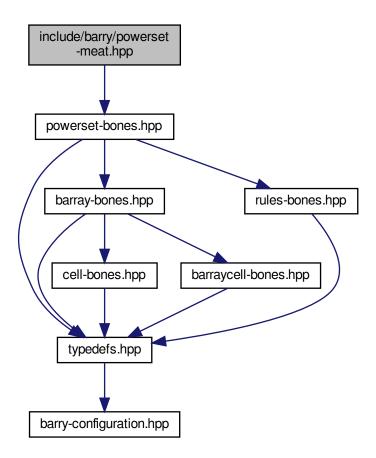


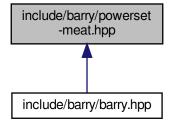
# **Classes**

# 8.38 include/barry/powerset-meat.hpp File Reference

#include "powerset-bones.hpp"

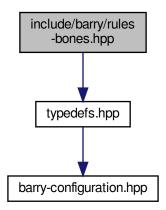
Include dependency graph for powerset-meat.hpp:



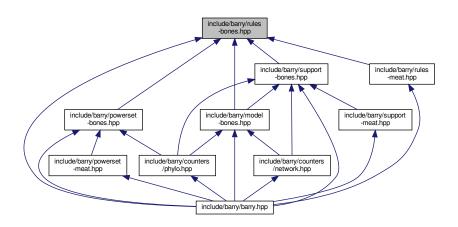


# 8.39 include/barry/rules-bones.hpp File Reference

#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.
- 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)

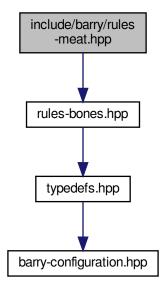
#### 8.39.1 Function Documentation

#### 8.39.1.1 rule\_fun\_default()

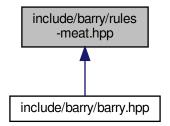
Definition at line 10 of file rules-bones.hpp.

# 8.40 include/barry/rules-meat.hpp File Reference

```
#include "rules-bones.hpp"
Include dependency graph for rules-meat.hpp:
```



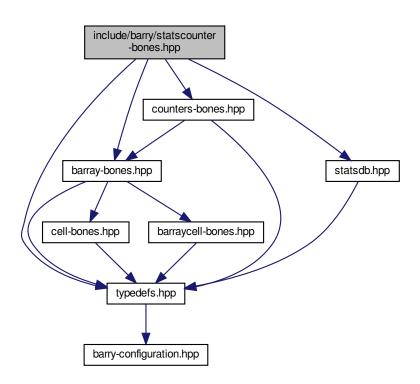
This graph shows which files directly or indirectly include this file:



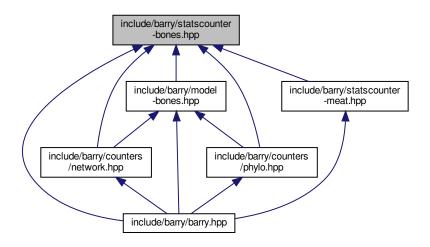
# 8.41 include/barry/statscounter-bones.hpp File Reference

```
#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:



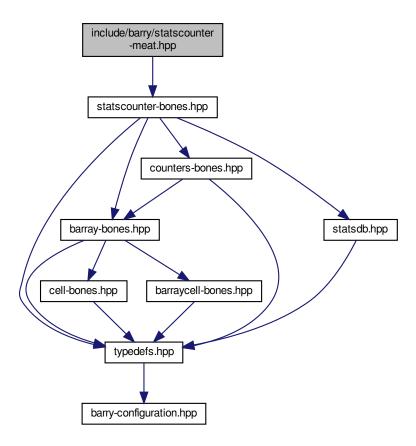
# Classes

class StatsCounter< Array\_Type, Data\_Type >
 Count stats for a single Array.

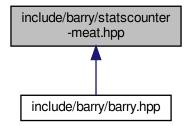
# 8.42 include/barry/statscounter-meat.hpp File Reference

#include "statscounter-bones.hpp"

Include dependency graph for statscounter-meat.hpp:



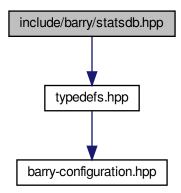
This graph shows which files directly or indirectly include this file:



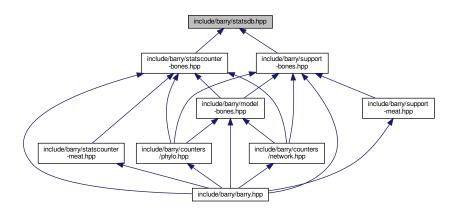
# 8.43 include/barry/statsdb.hpp File Reference

#include "typedefs.hpp"

Include dependency graph for statsdb.hpp:



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



#### **Classes**

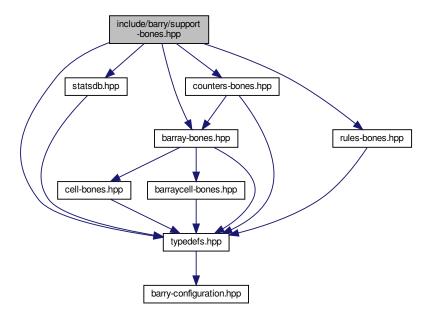
class FreqTable < T >

Database of statistics.

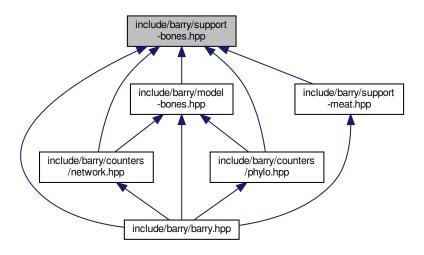
# 8.44 include/barry/support-bones.hpp File Reference

```
#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:

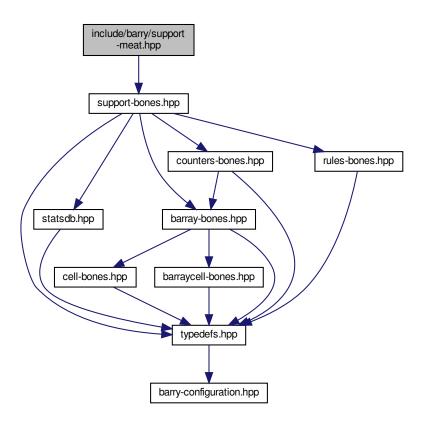


# **Classes**

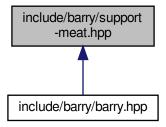
class Support < Array\_Type, Data\_Counter\_Type, Data\_Rule\_Type, Data\_Rule\_Dyn\_Type >
 Compute the support of sufficient statistics.

# 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</li>
- 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

```
 \begin{tabular}{ll} \# define & SUPPORT\_TEMPLATE ( & & \\ & a, & \\ & b & ) \\ \end{tabular}
```

#### 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()

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()

```
EmptyArray insert_cell (
          cfree. first,
          cfree. second,
          EmptyArray.default_val(). value,
          false ,
          false )
```

# 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]

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]

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

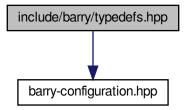
```
std::vector< Array_Type > std::vector< std::vector< double > >* stats_bank

Initial value:
{
    if (pos >= coordinates_free.size())
        return
```

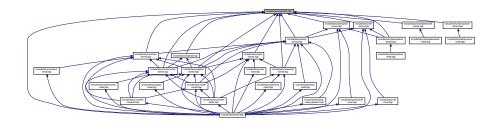
Definition at line 110 of file support-meat.hpp.

# 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

# **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 >

T vec_inner_prod (const std::vector< T > &a, const std::vector< T > &b)
```

```
    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::UKNOWN = -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 >, 151
    BArray< Cell Type, Data Type >, 31
\simBArrayCell
                                                      \simStatsCounter
                                                           StatsCounter< Array_Type, Data_Type >, 155
    BArrayCell< Cell_Type, Data_Type >, 42
~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 >, 155
\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 >, 144
\simGeese
                                                          Rules < Array_Type, Data_Type >, 152
    Geese, 105
                                                          Support<
                                                                       Array Type,
                                                                                      Data Counter Type,
\simModel
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
    Model<
                Array Type,
                                Data Counter Type,
                                                               161
         Data_Rule_Type, Data_Rule_Dyn_Type >,
                                                      add rule dyn
         117
                                                          Model <
                                                                      Array_Type,
                                                                                       Data Counter Type,
\simNetCounterData
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
    NetCounterData, 127
                                                               119, 120
\simNetworkData
                                                           Support<
                                                                       Array_Type,
                                                                                       Data_Counter_Type,
    NetworkData, 129
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
\simNode
                                                               162
    Node, 132
                                                      annotations
\simPhyloRuleDynData
                                                          Node, 133
    PhyloRuleDynData, 140
                                                      Array
\simPowerSet
                                                           ConstBArrayRowIter< Cell Type, Data Type >, 80
    PowerSet < Array_Type, Data_Rule_Type >, 144
                                                      arrav
\simRule
                                                          Node, 133
     Rule < Array_Type, Data_Type >, 149
                                                      array_bank
```

support-meat.hpp, 232	BARRY_BARRAY_MEAT_OPERATORS_HPP,
arrays	172
Node, 133	checkdim_, 172
AS_ONE	COL, 172
EXISTS, 25	ROW, 172
as_vector	barray-meat.hpp
FreqTable < T >, 100	COL, 174
AS_ZERO	ROW, 174
EXISTS, 25	BARRAY_BONES_HPP
at PhyloCounterDate 120	barray-bones.hpp, 170
PhyloCounterData, 138	BArrayCell Cell Type Data Type > 41
BArray	BArrayCell< Cell_Type, Data_Type >, 41 BArrayCell< Cell_Type, Data_Type >, 41
BArray< Cell_Type, Data_Type >, 30, 31	~BArrayCell, 42
BArray< Cell_Type, Data_Type >, 27	-
~BArray, 31	BArray Cell_Type, Data_Type >, 40
BArray, 30, 31	BArray Danas Call Type Data Type > 61
BArrayCell< Cell_Type, Data_Type >, 40	BArrayDense < Cell_Type, Data_Type >, 61
BArrayCell_const< Cell_Type, Data_Type >, 40	operator Cell_Type, 42
clear, 31	operator*=, 42
col, 31	operator+=, 42
D, 32	operator-=, 42
default_val, 32	operator/=, 43
flush_data, 32	operator=, 43
get_cell, 32	operator==, 43
get_col_vec, 32, 33	BArrayCell_const
get_entries, 33	BArrayCell_const < Cell_Type, Data_Type >, 44
get_row_vec, 33	BArrayCell_const < Cell_Type, Data_Type >, 43
insert_cell, 33, 34	~BArrayCell_const, 44
is_empty, 34	BArray Cell_Type, Data_Type >, 40
ncol, 34	BArray Dance & Call Time Data Time > 61
nnozero, 34	BArrayDense < Cell_Type, Data_Type >, 61
nrow, 35	operator Cell_Type, 44
operator*=, 35	operator!=, 45
operator(), 35	operator < .45
operator+=, 35, 36	operator 45
operator-=, 36	operator>, 45
operator/=, 36	operator>=, 46
operator=, 36, 37	operator==, 45
operator==, 37	BArray Danas Call Type Data Type > 40, 50
out_of_range, 37	BArrayDense < Cell_Type, Data_Type >, 49, 50 BArrayDense < Cell_Type, Data_Type >, 46
print, 37	~BArrayDense, 50
reserve, 37	-
resize, 37	BArrayCell< Cell_Type, Data_Type >, 61 BArrayCell_const< Cell_Type, Data_Type >, 61
rm_cell, 38	BArrayDense, 49, 50
row, 38	clear, 50
set_data, 38	col, 51
swap cells, 38	
swap_cols, 39	D, 51
swap_rows, 39	default_val, 51
toggle_cell, 39	get_cell, 51
toggle_lock, 39	get_col_vec, 52 get_entries, 52
transpose, 39	get_rentries, 52 get_row_vec, 52, 53
visited, 40	
zero_col, 40	insert_cell, 53
zero row, 40	is_empty, 54
barray-bones.hpp	ncol, 54
BARRAY_BONES_HPP, 170	nnozero, 54
barray-meat-operators.hpp	nrow, 54

operator*=, 55 operator(), 54, 55 operator+=, 55 operator-=, 56 operator/=, 56 operator==, 56, 57 operator==, 57 out_of_range, 57 print, 57 reserve, 57 resize, 58 rm_cell, 58 row, 58 set_data, 58 swap_cells, 59 swap_cols, 59	BArrayVector< Cell_Type, Data_Type >, 67  BArrayVector< Cell_Type, Data_Type >, 67  ~BArrayVector, 68  BArrayVector, 67  begin, 68  end, 68  is_col, 68  is_row, 69  operator std::vector< Cell_Type >, 69  operator+=, 69  operator+=, 69  operator-=, 69  operator-=, 70  operator==, 70  size, 70
swap_rows, 59	barrayvector-meat.hpp
toggle_cell, 59	BARRY_BARRAYVECTOR_MEAT_HPP, 183
toggle_lock, 60	BArrayVector_const
transpose, 60 visited, 61	BArrayVector_const< Cell_Type, Data_Type >, 71 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, 178	end, 72
COL, 178	is_col, 72
POS, 179	is_row, 72
ROW, 179	operator std::vector< Cell_Type >, 72
ZERO_CELL, 179	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 barry, 23
operator*=, 63 operator+=, 63	barry-configuration.hpp
operator-=, 63	BARRY_CHECK_SUPPORT, 184
operator/=, 63	BARRY_ISFINITE, 184
operator=, 63	BARRY_MAX_NUM_ELEMENTS, 184
operator==, 64	BARRY_SAFE_EXP, 184
barraydensecell-meat.hpp	Map, 184
BARRY_BARRAYDENSECELL_MEAT_HPP, 181	printf_barry, 184
POS, 181	barry.hpp
BArrayDenseCell_const	BARRY_HPP, 186
BArrayDenseCell_const< Cell_Type, Data_Type	BARRY_VERSION, 186
>, 65	COUNTER_FUNCTION, 186
BArrayDenseCell_const< Cell_Type, Data_Type >, 64	COUNTER_LAMBDA, 186
~BArrayDenseCell_const, 65	RULE_FUNCTION, 187
BArrayDenseCell_const, 65	RULE_LAMBDA, 187
operator Cell_Type, 65	barry::counters, 23
operator < 65	barry::counters::network, 24 barry::counters::phylo, 24
operator<, 65 operator<=, 66	BARRY_BARRAY_MEAT_OPERATORS_HPP
operator>, 66	barray-meat-operators.hpp, 172
operator>=, 66	BARRY_BARRAYDENSE_MEAT_HPP
operator==, 66	barraydense-meet.hpp, 178
BArrayVector	BARRY_BARRAYDENSECELL_MEAT_HPP

barraydensecell-meat.hpp, 181	NONE, 24
BARRY_BARRAYVECTOR_MEAT_HPP	ONE, 24
barrayvector-meat.hpp, 183	TWO, 24
BARRY_CHECK_SUPPORT	checkdim_
barry-configuration.hpp, 184	barray-meat-operators.hpp, 172
BARRY_HPP	clear
barry.hpp, 186	BArray< Cell_Type, Data_Type >, 31
BARRY_ISFINITE	BArrayDense < Cell_Type, Data_Type >, 50
barry-configuration.hpp, 184	Counters < Array_Type, Data_Type >, 88
BARRY_MAX_NUM_ELEMENTS	FreqTable < T >, 100
barry-configuration.hpp, 184	Rules < Array_Type, Data_Type >, 152
BARRY_SAFE_EXP	COL
barry-configuration.hpp, 184	barray-meat-operators.hpp, 172
BARRY_SUPPORT_MEAT_HPP	barray-meat.hpp, 174
support-meat.hpp, 226	barraydense-meet.hpp, 178
BARRY_VERSION	col
barry.hpp, 186	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, 237
PhyloCounterData, 138	colnames
PowerSet < Array_Type, Data_Rule_Type >, 144	Flock, 94
blengths	Geese, 105
NodeData, 136	Model < Array_Type, Data_Counter_Type,
BOTH	Data_Rule_Type, Data_Rule_Dyn_Type >,
CHECK, 24	120
EXISTS, 25	ConstBArrayRowlter
calc	ConstBArrayRowlter< Cell_Type, Data_Type >, 79
PowerSet< Array_Type, Data_Rule_Type >, 145	ConstBArrayRowlter< Cell_Type, Data_Type >, 79
Support< Array_Type, Data_Counter_Type,	~ConstBArrayRowIter, 80
Data_Rule_Type, Data_Rule_Dyn_Type >,	Array, 80
162	ConstBArrayRowlter, 79
calc backend	current_col, 80
support-meat.hpp, 227	current_row, 80
calc_reduced_sequence	iter, 80
Geese, 105	coordinates_free
calc_sequence	PowerSet < Array_Type, Data_Rule_Type >, 146
Geese, 105	Support Array_Type, Data_Counter_Type,
Cell	Data_Rule_Type, Data_Rule_Dyn_Type >,
Cell< Cell_Type >, 75, 76	166
Cell< Cell_Type >, 74	coordinates_locked PowerSet< Array Type, Data Rule Type >, 146
~Cell, 75	· · · · · · · · · · · · · · ·
add, 76, 77	
Cell, 75, 76	Data_Rule_Type, Data_Rule_Dyn_Type >, 166
operator Cell_Type, 77	
operator!=, 77	Country CArroy Type Date Type > 02
operator=, 77, 78	Counter< Array_Type, Data_Type >, 83
operator==, 78	count_all
value, 78	StatsCounter< Array_Type, Data_Type >, 156
visited, 78	count_current  StateCounter < Array Type Data Type > 156
cfree	StatsCounter< Array_Type, Data_Type >, 156
support-meat.hpp, 232	count_fun
change_stats	Counter< Array_Type, Data_Type >, 84
Support< Array_Type, Data_Counter_Type,	count_init
Data_Rule_Type, Data_Rule_Dyn_Type >,	StatsCounter< Array_Type, Data_Type >, 156
165	Counter Array Type Data Type > 92 92
CHECK, 24	Counter < Array Type, Data Type >, 82, 83
BOTH, 24	Counter< Array_Type, Data_Type >, 81
•	

$\sim$ Counter, 83	Phylo counters, 18
count, 83	counter_nodecov
count_fun, 84	Network counters, 13
Counter, 82, 83	counter_nodeicov
data, 84	Network counters, 13
delete_data, 85	counter_nodematch
desc, 85	Network counters, 13
init, 83	counter_nodeocov
init_fun, 85	Network counters, 13
name, 85	counter_odegree
operator=, 84	Network counters, 14
counter_absdiff	counter_odegree15
Network counters, 11	Network counters, 14
counter_co_opt	counter_ostar2
Phylo counters, 16	Network counters, 14
counter_cogain	counter_overall_changes
Phylo counters, 16	Phylo counters, 18
counter_ctriads	counter_overall_gains
Network counters, 11	Phylo counters, 19
counter_degree	counter_overall_loss
Network counters, 11	Phylo counters, 19
counter_density	counter_prop_genes_changing
Network counters, 11	Phylo counters, 19
counter_diff	counter_subfun
Network counters, 11	Phylo counters, 19
counter_edges	counter_ttriads
Network counters, 12	Network counters, 14
Counter_fun_type	Counters
typedefs.hpp, 237	Counters < Array_Type, Data_Type >, 86, 87
COUNTER FUNCTION	counters
barry.hpp, 186	support-meat.hpp, 232
counter_gains	Counters< Array_Type, Data_Type >, 86
Phylo counters, 16	~Counters, 87
counter_gains_k_offspring	add_counter, 88
Phylo counters, 17	clear, 88
counter genes changing	Counters, 86, 87
Phylo counters, 17	operator=, 88, 89
counter_idegree	operator[], 89
Network counters, 12	size, 90
counter_idegree15	counters
Network counters, 12	support-meat.hpp, 232
counter_isolates	Counting, 9
Network counters, 12	counts
counter_istar2	PhyloRuleDynData, 141
Network counters, 12	Counts_type
COUNTER_LAMBDA	typedefs.hpp, 237
barry.hpp, 186	current col
counter_longest	ConstBArrayRowIter< Cell Type, Data Type >, 80
Phylo counters, 17	current_row
counter_loss	ConstBArrayRowIter< Cell_Type, Data_Type >, 80
Phylo counters, 17	current_stats
counter_maxfuns	Support< Array_Type, Data_Counter_Type
Phylo counters, 18	Data_Rule_Type, Data_Rule_Dyn_Type >
counter_mutual	166
Network counters, 13	
counter_neofun	D
Phylo counters, 18	BArray< Cell_Type, Data_Type >, 32
counter_neofun_a2b	BArrayDense< Cell_Type, Data_Type >, 51
555to:_!!65!d!!_dE5	Rule < Array_Type, Data_Type >, 150

dat	Support< Array_Type, Data_Counter_Type,
Flock, 98	Data_Rule_Type, Data_Rule_Dyn_Type >
data	162
Counter< Array_Type, Data_Type >, 84	EXISTS, 25
PowerSet < Array_Type, Data_Rule_Type >, 147	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
166	
support-meat.hpp, 233	f_
delete_data	support-meat.hpp, 233
Counter< Array_Type, Data_Type >, 85	Flock, 92
delete_rengine	$\sim$ Flock, 94
Geese, 112	add_data, 94
delete_rules	colnames, 94
Support< Array_Type, Data_Counter_Type,	dat, 98
Data_Rule_Type, Data_Rule_Dyn_Type >,	Flock, 93
166	get_counters, 94
support-meat.hpp, 233	get_model, 95
delete_rules_dyn	get_support, 95
Support< Array_Type, Data_Counter_Type,	init, 95
Data_Rule_Type, Data_Rule_Dyn_Type >,	initialized, 98
167	likelihood_joint, 95
support-meat.hpp, 233	model, 98
delete_support	nfunctions, 98
Geese, 112	nfuns, 96
desc	nleafs, 96
Counter< Array_Type, Data_Type >, 85	nnodes, 96
directed	nterms, 96
NetworkData, 129	ntrees, 96
duplication	operator(), 96
Node, 134	parse_polytomies, 97
NodeData, 137	print, 97
PhyloRuleDynData, 141	rengine, 98
	set_seed, 97
else	support_size, 97
support-meat.hpp, 233	flush_data
empty	BArray< Cell_Type, Data_Type >, 32
PhyloCounterData, 138	for
EmptyArray	support-meat.hpp, 228
PowerSet < Array_Type, Data_Rule_Type >, 147	FreqTable
end	FreqTable < T >, 99
BArrayVector< Cell_Type, Data_Type >, 68	FreqTable < T >, 99
BArrayVector_const< Cell_Type, Data_Type >, 72	∼FreqTable, 99
PhyloCounterData, 138	add, 100
PowerSet < Array_Type, Data_Rule_Type >, 145	as_vector, 100
Entries Coll. Time > 04	clear, 100
Entries < Cell_Type >, 91	FreqTable, 99
Entries < Cell_Type >, 90	get_data, 100
~Entries, 91	get_data_ptr, 100
Entries, 91	print, 100
resize, 91	reserve, 101
source, 92	size, 101
target, 92	Conn. 101
val, 92	Geese, 101
eval_rules_dyn	$\sim$ Geese, 105

calc_reduced_sequence, 105	get_col_vec
calc_sequence, 105	BArray< Cell_Type, Data_Type >, 32, 33
colnames, 105	BArrayDense < Cell_Type, Data_Type >, 52
delete_rengine, 112	get_counters
delete_support, 112	Flock, 94
Geese, 104	Geese, 106
get_annotated_nodes, 105	Model < Array_Type, Data_Counter_Type,
get_counters, 106	Data_Rule_Type, Data_Rule_Dyn_Type >,
get_model, 106	120
get_probabilities, 106	PhyloCounterData, 139
get_rengine, 106	StatsCounter< Array_Type, Data_Type >, 156
get_states, 106	Support< Array_Type, Data_Counter_Type,
get_support, 107	Data_Rule_Type, Data_Rule_Dyn_Type >,
inherit_support, 107	163
init, 107	get_counts
init_node, 107	Support< Array_Type, Data_Counter_Type,
initialized, 112	Data_Rule_Type, Data_Rule_Dyn_Type >,
likelihood, 107	163
likelihood_exhaust, 108	get_counts_ptr
map_to_nodes, 112	Support< Array_Type, Data_Counter_Type,
nannotations, 108	Data_Rule_Type, Data_Rule_Dyn_Type >,
nfunctions, 112	163
nfuns, 108	get_current_stats
nleafs, 108	Support< Array_Type, Data_Counter_Type,
nnodes, 108	Data_Rule_Type, Data_Rule_Dyn_Type >,
nodes, 113	163
nterms, 109	get_data
observed_counts, 109	FreqTable $<$ T $>$ , 100
operator=, 109	PowerSet < Array_Type, Data_Rule_Type >, 145
parse_polytomies, 109	Support< Array_Type, Data_Counter_Type,
predict, 109	Data_Rule_Type, Data_Rule_Dyn_Type >,
predict_backend, 110	163
predict_exhaust, 110	get_data_ptr
predict_exhaust_backend, 110	FreqTable $<$ T $>$ , 100
predict_sim, 110	PowerSet < Array_Type, Data_Rule_Type >, 145
print, 110	get_entries
print_observed_counts, 111	BArray< Cell_Type, Data_Type >, 33
reduced_sequence, 113	BArrayDense < Cell_Type, Data_Type >, 52
sequence, 113	get_last_name
set_seed, 111	phylo.hpp, 202
simulate, 111	get_model
support_size, 111	Flock, 95
update_annotations, 111	Geese, 106
geese-bones.hpp	get_norm_const
INITIALIZED, 208	Model < Array_Type, Data_Counter_Type,
keygen_full, 209	Data_Rule_Type, Data_Rule_Dyn_Type >,
RULE_FUNCTION, 209	120
vec_diff, 209	get_parent
vector_caster, 209	Node, 132
gen_key	get_probabilities
Model< Array_Type, Data_Counter_Type,	Geese, 106
Data_Rule_Type, Data_Rule_Dyn_Type >,	get_pset
120	Model Array_Type, Data_Counter_Type,
get_annotated_nodes	Data_Rule_Type, Data_Rule_Dyn_Type >,
Geese, 105	121
get_cell	get_pset_stats
BArray Cell_Type, Data_Type >, 32	Model < Array_Type, Data_Counter_Type,
BArrayDense< Cell_Type, Data_Type >, 51	Data_Rule_Type, Data_Rule_Dyn_Type >,

121	include/barry/model-bones.hpp, 203
get_rengine	include/barry/model-meat.hpp, 204
Geese, 106	include/barry/models/geese.hpp, 206
Model Array_Type, Data_Counter_Type,	include/barry/models/geese/flock-bones.hpp, 207
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/models/geese/flock-meet.hpp, 207
121	include/barry/models/geese/geese-bones.hpp, 208
get row vec	include/barry/models/geese/geese-meat-constructors.hpp,
BArray< Cell_Type, Data_Type >, 33	210
BArrayDense < Cell_Type, Data_Type >, 52, 53	include/barry/models/geese/geese-meat-likelihood.hpp,
get_rules	210
Model< Array_Type, Data_Counter_Type,	include/barry/models/geese/geese-meat-likelihood_exhaust.hpp,
Data_Rule_Type, Data_Rule_Dyn_Type >,	211
121	include/barry/models/geese/geese-meat-predict.hpp,
Support	212
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/models/geese/geese-meat-predict_exhaust.hpp,
164	212
get_rules_dyn	include/barry/models/geese/geese-meat-predict_sim.hpp,
Model Array_Type, Data_Counter_Type,	213
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/models/geese/geese-meat-simulate.hpp,
121	213
Support< Array_Type, Data_Counter_Type,	include/barry/models/geese/geese-meat.hpp, 214
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/models/geese/geese-node-bones.hpp,
164	214
get_seq	include/barry/powerset-bones.hpp, 215
Rules < Array_Type, Data_Type >, 152	include/barry/powerset-meat.hpp, 216
get_states	include/barry/rules-bones.hpp, 218
Geese, 106	include/barry/rules-meat.hpp, 219
get_support	include/barry/statscounter-bones.hpp, 220
Flock, 95	include/barry/statscounter-meat.hpp, 221
Geese, 107	include/barry/statsdb.hpp, 222
Model Array_Type, Data_Counter_Type,	include/barry/support-bones.hpp, 223
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/support-meat.hpp, 225
122	include/barry/typedefs.hpp, 235
i.d	indices
Node, 134	NetCounterData, 127
if	inherit_support
	Geese, 107
support-meat.hpp, 228	init
include/barry/barray-bones.hpp, 169	Counter< Array_Type, Data_Type >, 83
include/barry/barray-iterator.hpp, 170	Flock, 95
include/barry/barray-meat-operators.hpp, 171	Geese, 107
include/barry/barray-meat.hpp, 173	init_fun
include/barry/barraycell-bones.hpp, 174	Counter< Array_Type, Data_Type >, 85
include/barry/barraydenee hance hanc	init_node
include/barry/barraydense-bones.hpp, 176	Geese, 107
include/barry/barraydense-meet.hpp, 177	init_support
include/barry/barraydensecell-bones.hpp, 179	PowerSet < Array_Type, Data_Rule_Type >, 145
include/barry/barraydensecell-meat.hpp, 180	Support< Array_Type, Data_Counter_Type,
include/barry/barrayvector-bones.hpp, 181	Data_Rule_Type, Data_Rule_Dyn_Type >,
include/barry/barrayvector-meat.hpp, 182	164
include/barry/barry-configuration.hpp, 183	INITIALIZED
include/barry/barry.hpp, 185	geese-bones.hpp, 208
include/barry/cell-bones.hpp, 187	initialized
include/barry/cell-meat.hpp, 188	Flock, 98
include/barry/col-bones.hpp, 189	Geese, 112
include/barry/counters-bones.hpp, 189	insert_cell
include/barry/counters-meat.hpp, 190	BArray< Cell_Type, Data_Type >, 33, 34
include/barry/counters/network.hpp, 191	BArrayDense < Cell_Type, Data_Type >, 53
include/barry/counters/phylo.hpp, 197	••

support-meat.hpp, 228	Model < Array_Type, Data_Counter_Type, Data_Rule_Type,
is_col	Data_Rule_Dyn_Type >, 113
BArrayVector< Cell_Type, Data_Type >, 68	$\sim$ Model, 117
BArrayVector_const< Cell_Type, Data_Type >, 72	add_array, 118
is_empty	add_counter, 118
BArray< Cell_Type, Data_Type >, 34	add_rule, 119
BArrayDense< Cell_Type, Data_Type >, 54	add_rule_dyn, 119, 120
is_leaf	colnames, 120
Node, 133	gen_key, 120
is_row	get_counters, 120
BArrayVector< Cell_Type, Data_Type >, 69	get_norm_const, 120
BArrayVector_const< Cell_Type, Data_Type >, 72	get_pset, 121
iter	get_pset_stats, 121
ConstBArrayRowlter< Cell_Type, Data_Type >, 80	get_rengine, 121
	get_rules, 121
keygen_default	get_rules_dyn, 121
model-bones.hpp, 204	get_support, 122
keygen_full	likelihood, 122
geese-bones.hpp, 209	likelihood total, 122
	Model, 117
lb	nterms, 123
PhyloRuleDynData, 141	operator=, 123
likelihood	print, 123
Geese, 107	print_stats, 123
Model< Array_Type, Data_Counter_Type,	sample, 123, 124
Data_Rule_Type, Data_Rule_Dyn_Type >,	
122	set_counters, 124
likelihood	set_keygen, 124
model-meat.hpp, 205	set_rengine, 124
likelihood_exhaust	set_rules, 124
Geese, 108	set_rules_dyn, 125
likelihood_joint	set_seed, 125
Flock, 95	size, 125
	size_unique, 125
likelihood_total  Model< Array Type, Data Counter Type,	store_psets, 125
3=31 / = = 31 /	support_size, 126
Data_Rule_Type, Data_Rule_Dyn_Type >,	model-bones.hpp
122	keygen_default, 204
M	model-meat.hpp
PowerSet< Array_Type, Data_Rule_Type >, 147	likelihood_, 205
Support< Array_Type, Data_Counter_Type,	MODEL_TEMPLATE, 205, 206
Data_Rule_Type, Data_Bule_Dyn_Type >,	MODEL_TEMPLATE_ARGS, 205
167	MODEL_TYPE, 205
	update_normalizing_constant, 206
Map	MODEL_TEMPLATE
barry-configuration.hpp, 184	model-meat.hpp, 205, 206
map_to_nodes	MODEL_TEMPLATE_ARGS
Geese, 112	model-meat.hpp, 205
MapVec_type	MODEL TYPE
typedefs.hpp, 237	model-meat.hpp, 205
max_num_elements	
Support< Array_Type, Data_Counter_Type,	N
Data_Rule_Type, Data_Rule_Dyn_Type >,	PowerSet < Array_Type, Data_Rule_Type >, 147
167	Support< Array_Type, Data_Counter_Type,
Model	Data_Rule_Type, Data_Rule_Dyn_Type >,
Model < Array_Type, Data_Counter_Type,	167
Data_Rule_Type, Data_Rule_Dyn_Type >,	name
117	Counter< Array_Type, Data_Type >, 85
model	nannotations
Flock, 98	Geese, 108
	_,,

narray	NetRules, 196
Node, 134	NetStatsCounter, 196
ncol	NetSupport, 196
BArray< Cell_Type, Data_Type >, 34	Network, 196
BArrayDense< Cell_Type, Data_Type >, 54	NETWORK_COUNTER, 194
NET_C_DATA_IDX	NETWORK_COUNTER_LAMBDA, 194
network.hpp, 194	NETWORK_RULE, 195
NET_C_DATA_NUM	NETWORK_RULE_LAMBDA, 195
network.hpp, 194	rules_zerodiag, 197
NetCounter	NETWORK COUNTER
network.hpp, 195	Network counters, 14
NetCounterData, 126	network.hpp, 194
$\sim$ NetCounterData, 127	NETWORK_COUNTER_LAMBDA
indices, 127	network.hpp, 194
NetCounterData, 126, 127	NETWORK RULE
numbers, 127	network.hpp, 195
NetCounters	NETWORK RULE LAMBDA
network.hpp, 195	network.hpp, 195
NetModel	117
	NetworkData, 128
network.hpp, 196	~NetworkData, 129
NetRule	directed, 129
network.hpp, 196	NetworkData, 128, 129
NetRules	vertex_attr, 129
network.hpp, 196	nfunctions
NetStatsCounter	Flock, 98
network.hpp, 196	Geese, 112
NetSupport	nfuns
network.hpp, 196	Flock, 96
Network	Geese, 108
network.hpp, 196	nleafs
Network counters, 10	Flock, 96
counter_absdiff, 11	Geese, 108
counter_ctriads, 11	nnodes
counter_degree, 11	Flock, 96
counter_density, 11	Geese, 108
counter_diff, 11	nnozero
counter edges, 12	BArray< Cell_Type, Data_Type >, 34
counter idegree, 12	BArrayDense< Cell_Type, Data_Type >, 54
counter_idegree15, 12	Node, 130
counter isolates, 12	~Node, 132
counter_istar2, 12	annotations, 133
counter mutual, 13	array, 133
counter_nodecov, 13	arrays, 133
counter_nodeicov, 13	duplication, 134
counter_nodematch, 13	get_parent, 132
counter nodeocov, 13	id, 134
counter_nodeocov, 13	
_ •	
	is_leaf, 133
counter_odegree15, 14	narray, 134
counter_ostar2, 14	narray, 134 Node, 131, 132
counter_ostar2, 14 counter_ttriads, 14	narray, 134 Node, 131, 132 noffspring, 133
counter_ostar2, 14 counter_ttriads, 14 NETWORK_COUNTER, 14	narray, 134 Node, 131, 132 noffspring, 133 offspring, 134
counter_ostar2, 14 counter_ttriads, 14 NETWORK_COUNTER, 14 network.hpp	narray, 134 Node, 131, 132 noffspring, 133 offspring, 134 ord, 134
counter_ostar2, 14 counter_ttriads, 14 NETWORK_COUNTER, 14 network.hpp NET_C_DATA_IDX, 194	narray, 134 Node, 131, 132 noffspring, 133 offspring, 134 ord, 134 parent, 135
counter_ostar2, 14 counter_ttriads, 14 NETWORK_COUNTER, 14 network.hpp NET_C_DATA_IDX, 194 NET_C_DATA_NUM, 194	narray, 134 Node, 131, 132 noffspring, 133 offspring, 134 ord, 134 parent, 135 probability, 135
counter_ostar2, 14 counter_ttriads, 14 NETWORK_COUNTER, 14 network.hpp NET_C_DATA_IDX, 194 NET_C_DATA_NUM, 194 NetCounter, 195	narray, 134 Node, 131, 132 noffspring, 133 offspring, 134 ord, 134 parent, 135 probability, 135 subtree_prob, 135
counter_ostar2, 14 counter_ttriads, 14 NETWORK_COUNTER, 14 network.hpp NET_C_DATA_IDX, 194 NET_C_DATA_NUM, 194 NetCounter, 195 NetCounters, 195	narray, 134 Node, 131, 132 noffspring, 133 offspring, 134 ord, 134 parent, 135 probability, 135 subtree_prob, 135 visited, 135
counter_ostar2, 14 counter_ttriads, 14 NETWORK_COUNTER, 14 network.hpp NET_C_DATA_IDX, 194 NET_C_DATA_NUM, 194 NetCounter, 195 NetCounters, 195 NetModel, 196	narray, 134 Node, 131, 132 noffspring, 133 offspring, 134 ord, 134 parent, 135 probability, 135 subtree_prob, 135 visited, 135 NodeData, 136
counter_ostar2, 14 counter_ttriads, 14 NETWORK_COUNTER, 14 network.hpp NET_C_DATA_IDX, 194 NET_C_DATA_NUM, 194 NetCounter, 195 NetCounters, 195	narray, 134 Node, 131, 132 noffspring, 133 offspring, 134 ord, 134 parent, 135 probability, 135 subtree_prob, 135 visited, 135

duplication, 137	BArrayDenseCell_const< Cell_Type, Data_Type
NodeData, 136	>, 66
states, 137	BArrayVector_const< Cell_Type, Data_Type >, 73
nodes	operator>=  PArroyColl const < Coll Type Data Type > 46
Geese, 113	BArrayCell_const < Cell_Type, Data_Type >, 46
noffspring Node, 133	BArrayDenseCell_const< Cell_Type, Data_Type >, 66
NONE	•
	BArrayVector_const< Cell_Type, Data_Type >, 73
CHECK, 24 EXISTS, 26	operator*= BArray< Cell_Type, Data_Type >, 35
NYOW  PArroy Coll Type Date Type > 25	BArrayCell < Cell_Type, Data_Type >, 42
BArray Danas Coll Time 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, 139
ntrees	Rule < Array_Type, Data_Type >, 150
Flock, 96	Rules < Array_Type, Data_Type >, 153
numbers	vecHasher< T >, 168
NetCounterData, 127	operator+=
observed_counts	BArray< Cell_Type, Data_Type >, 35, 36
Geese, 109	BArrayCell< Cell_Type, Data_Type >, 42
offspring	BArrayDense < Cell_Type, Data_Type >, 55
Node, 134	BArrayDenseCell Cell_Type, Data_Type >, 63
ONE	BArrayVector< Cell_Type, Data_Type >, 69
CHECK, 24	operator-=
EXISTS, 26	BArray< Cell_Type, Data_Type >, 36
operator Cell_Type	BArrayCell< Cell_Type, Data_Type >, 42
BArrayCell < Cell_Type, Data_Type >, 42	BArrayDense < Cell_Type, Data_Type >, 56
BArrayCell_const< Cell_Type, Data_Type >, 44	BArrayDenseCell< Cell_Type, Data_Type >, 63
BArrayDenseCell< Cell_Type, Data_Type >, 44  BArrayDenseCell< Cell_Type, Data_Type >, 62	BArrayVector< Cell_Type, Data_Type >, 69
BArrayDenseCell_const< Cell_Type, Data_Type  BarrayDenseCell_const< Cell_Type, Data_Type	operator/=
>, 65	BArray< Cell_Type, Data_Type >, 36
Cell< Cell_Type >, 77	BArrayCell< Cell_Type, Data_Type >, 43
operator std::vector< Cell_Type >	BArrayDense < Cell_Type, Data_Type >, 56
BArrayVector < Cell_Type, Data_Type >, 69	BArrayDenseCell< Cell_Type, Data_Type >, 63
BArrayVector_const< Cell_Type, Data_Type >, 72	BArrayVector< Cell_Type, Data_Type >, 70
operator!=	operator=
BArrayCell_const< Cell_Type, Data_Type >, 45	BArray< Cell_Type, Data_Type >, 36, 37
BArrayDenseCell_const< Cell_Type, Data_Type  BarrayDenseCell_const< Cell_Type, Data_Type	BArrayCell< Cell_Type, Data_Type >, 43
>, 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
BArrayCell_const< Cell_Type, Data_Type >, 45	Counter< Array_Type, Data_Type >, 84
BArrayDenseCell_const< Cell_Type, Data_Type  BarrayDenseCell_const< Cell_Type, Data_Type	Counters < Array_Type, Data_Type >, 88, 89
>, 65	Geese, 109
BArrayVector_const< Cell_Type, Data_Type >, 73	Model< Array_Type, Data_Counter_Type,
operator<=	Data_Rule_Type, Data_Rule_Dyn_Type >,
BArrayCell_const< Cell_Type, Data_Type >, 45	123
BArrayDenseCell_const< Cell_Type, Data_Type  BArrayDenseCell_const< Cell_Type, Data_Type	Rules < Array_Type, Data_Type >, 153
>, 66	operator==
	BArray< Cell_Type, Data_Type >, 37
BArrayVector_const< Cell_Type, Data_Type >, 73	BArrayCell< Cell_Type, Data_Type >, 43
operator>  RArrayCall const < Call Type Data Type > 45	BArrayCell_const< Cell_Type, Data_Type >, 45
BArrayCell_const< Cell_Type, Data_Type >, 45	

BArrayDense< Cell_Type, Data_Type >, 57 BArrayDenseCell< Cell_Type, Data_Type >, 64	phylo.hpp, 199 PHYLO_RULE_DYN_LAMBDA
BArrayDenseCell_const< Cell_Type, Data_Type	phylo.hpp, 200
>, 66	PhyloArray
BArrayVector< Cell_Type, Data_Type >, 70	phylo.hpp, 200
BArrayVector_const< Cell_Type, Data_Type >, 73	PhyloCounter
Cell< Cell_Type >, 78	phylo.hpp, 200
operator[]	PhyloCounterData, 137
Counters< Array_Type, Data_Type >, 89	at, 138
PowerSet < Array_Type, Data_Rule_Type >, 146	begin, 138
ord	empty, 138
Node, 134	end, 138
out_of_range	get_counters, 139
BArray < Cell_Type, Data_Type >, 37	operator(), 139
BArrayDense< Cell_Type, Data_Type >, 57	PhyloCounterData, 138
parent	push_back, 139
Node, 135	reserve, 139
parse_polytomies	shrink_to_fit, 139
Flock, 97	size, 139
Geese, 109	PhyloCounters
Phylo counters, 15	phylo.hpp, 200 PhyloModel
counter_co_opt, 16	phylo.hpp, 201
counter_cogain, 16	PhyloPowerSet
counter_gains, 16	phylo.hpp, 201
counter_gains_k_offspring, 17	PhyloRule
counter_genes_changing, 17	phylo.hpp, 201
counter_longest, 17	PhyloRuleData
counter_loss, 17	phylo.hpp, 201
counter_maxfuns, 18	PhyloRuleDyn
counter_neofun, 18	phylo.hpp, 201
counter_neofun_a2b, 18	PhyloRuleDynData, 140
counter_overall_changes, 18	∼PhyloRuleDynData, 140
counter_overall_gains, 19	counts, 141
counter_overall_loss, 19	duplication, 141
counter_prop_genes_changing, 19	lb, 141
counter_subfun, 19	PhyloRuleDynData, 140
Phylo rules, 20	pos, 141
rule_dyn_limit_changes, 20	ub, 141
phylo.hpp	PhyloRules
get_last_name, 202	phylo.hpp, 201
PHYLO_CHECK_MISSING, 199	PhyloRulesDyn
PHYLO_COUNTER_LAMBDA, 199	phylo.hpp, 202
PHYLO_RULE_DYN_LAMBDA, 200	PhyloStatsCounter
PhyloArray, 200 PhyloCounter, 200	phylo.hpp, 202
PhyloCounters, 200 PhyloCounters, 200	PhyloSupport
PhyloModel, 201	phylo.hpp, 202
PhyloPowerSet, 201	POS
PhyloRule, 201	barraydense-meet.hpp, 179
PhyloRuleData, 201	barraydensecell-meat.hpp, 181
PhyloRuleDyn, 201	pos
PhyloRules, 201	PhyloRuleDynData, 141
PhyloRulesDyn, 202	PowerSet < Array Type Data Bula Type > 143
PhyloStatsCounter, 202	PowerSet < Array_Type, Data_Rule_Type >, 143
PhyloSupport, 202	PowerSet Array_Type, Data_Rule_Type >, 142
PHYLO_CHECK_MISSING	~PowerSet, 144 add_rule, 144
phylo.hpp, 199	begin, 144
PHYLO COUNTER LAMBDA	begin, 144

calc, 145	BArrayDense< Cell_Type, Data_Type >, 57
coordinates_free, 146 coordinates_locked, 146	FreqTable < T >, 101
<del>-</del> · · ·	PhyloCounterData, 139
data, 147	PowerSet < Array Type Data Bula Type > 146
EmptyArray, 147	PowerSet < Array_Type, Data_Rule_Type >, 146
end, 145	reset_array
get_data, 145	StatsCounter< Array_Type, Data_Type >, 156
get_data_ptr, 145	Support< Array_Type, Data_Counter_Type,
init_support, 145	Data_Rule_Type, Data_Rule_Dyn_Type >,
M, 147	164, 165
N, 147	resize  PArroy < Coll Type Data Type > 27
operator[], 146	BArray Cell_Type, Data_Type >, 37
PowerSet, 143	BArrayDense < Cell_Type, Data_Type >, 58
reset, 146	Entries < Cell_Type >, 91
rules, 147	return
rules_deleted, 148	support-meat.hpp, 234
size, 146	rm_cell
predict	BArray< Cell_Type, Data_Type >, 38
Geese, 109	BArrayDense< Cell_Type, Data_Type >, 58
predict_backend	support-meat.hpp, 228
Geese, 110	ROW
predict_exhaust	barray-meat-operators.hpp, 172
Geese, 110	barray-meat.hpp, 174
predict_exhaust_backend	barraydense-meet.hpp, 179
Geese, 110	row
predict_sim	BArray< Cell_Type, Data_Type >, 38
Geese, 110	BArrayDense < Cell_Type, Data_Type >, 58
print	Row_type
BArray< Cell_Type, Data_Type >, 37	typedefs.hpp, 237
BArrayDense< Cell_Type, Data_Type >, 57	Rule
Flock, 97	Rule < Array_Type, Data_Type >, 149
FreqTable < T >, 100	Rule< Array_Type, Data_Type >, 148
Geese, 110	$\sim$ Rule, 149
Model< Array_Type, Data_Counter_Type,	D, 150
Data_Rule_Type, Data_Rule_Dyn_Type >,	operator(), 150
123	Rule, 149
Support< Array_Type, Data_Counter_Type,	rule_dyn_limit_changes
Data_Rule_Type, Data_Rule_Dyn_Type >,	Phylo rules, 20
164	rule_fun_default
print_observed_counts	rules-bones.hpp, 219
Geese, 111	Rule_fun_type
print_stats	typedefs.hpp, 238
Model< Array_Type, Data_Counter_Type,	RULE_FUNCTION
Data_Rule_Type, Data_Rule_Dyn_Type >,	barry.hpp, 187
123	geese-bones.hpp, 209
printf_barry	RULE_LAMBDA
barry-configuration.hpp, 184	barry.hpp, 187
probability	Rules
Node, 135	Rules < Array_Type, Data_Type >, 151
push_back	rules
PhyloCounterData, 139	PowerSet < Array_Type, Data_Rule_Type >, 147
	support-meat.hpp, 234
README.md, 239	Rules< Array_Type, Data_Type >, 150
reduced_sequence	∼Rules, 151
Geese, 113	add_rule, 152
rengine	clear, 152
Flock, 98	get_seq, 152
reserve	operator(), 153
BArray< Cell_Type, Data_Type >, 37	• •

operator=, 153 Rules, 151 size, 153	shrink_to_fit PhyloCounterData, 139 simulate
rules-bones.hpp rule_fun_default, 219	Geese, 111 size
rules_ support-meat.hpp, 234	BArrayVector< Cell_Type, Data_Type >, 70 BArrayVector_const< Cell_Type, Data_Type >, 74
rules_deleted PowerSet< Array_Type, Data_Rule_Type >, 148	Counters < Array_Type, Data_Type >, 90 FreqTable < T >, 101
rules_dyn support-meat.hpp, 234	Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,
rules_zerodiag network.hpp, 197	125 PhyloCounterData, 139
sample	PowerSet< Array_Type, Data_Rule_Type >, 146 Rules< Array_Type, Data_Type >, 153
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 123, 124	size_unique  Model< Array_Type, Data_Counter_Type,  Data_Rule_Type, Data_Rule_Dyn_Type >,
sequence Geese, 113	125
set_counters	source Entries < Cell_Type >, 92
Model < Array_Type, Data_Counter_Type,	states
Data_Rule_Type, Data_Rule_Dyn_Type >, 124	NodeData, 137
StatsCounter< Array_Type, Data_Type >, 157	Statistical Models, 9 stats_bank
Support< Array_Type, Data_Counter_Type,	support-meat.hpp, 234
Data_Rule_Type, Data_Rule_Dyn_Type >,	StatsCounter
165	StatsCounter< Array_Type, Data_Type >, 155
set_data	StatsCounter< Array_Type, Data_Type >, 154
BArray< Cell_Type, Data_Type >, 38 BArrayDense< Cell_Type, Data_Type >, 58	~StatsCounter, 155
set_keygen	add_counter, 155
Model< Array_Type, Data_Counter_Type,	count_all, 156 count_current, 156
Data_Rule_Type, Data_Rule_Dyn_Type >,	count_init, 156
124	get_counters, 156
set_rengine	reset_array, 156
Model < Array_Type, Data_Counter_Type,	set_counters, 157
Data_Rule_Type, Data_Rule_Dyn_Type >, 124	StatsCounter, 155
set rules	store_psets  Madel < Array Type
	Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,
Data_Rule_Type, Data_Rule_Dyn_Type >,	125
124	subtree_prob
Support< Array_Type, Data_Counter_Type,	Node, 135
Data_Rule_Type, Data_Rule_Dyn_Type >, 165	Support
set_rules_dyn	Support< Array_Type, Data_Counter_Type,
Model< Array_Type, Data_Counter_Type,	Data_Rule_Type, Data_Rule_Dyn_Type >, 160
Data_Rule_Type, Data_Rule_Dyn_Type >,	Support < Array_Type, Data_Counter_Type, Data_Rule_Type
Support< Array_Type, Data_Counter_Type,	Data_Rule_Dyn_Type >, 157 ~Support, 161
Data_Rule_Type, Data_Rule_Dyn_Type >,	add_counter, 161
165	add_rule, 161
set_seed	add_rule_dyn, 162
Flock, 97	calc, 162
Geese, 111  Model< Array_Type, Data_Counter_Type,	change_stats, 165
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,	coordinates_free, 166
125	coordinates_locked, 166

current_stats, 166	swap_cells
delete_counters, 166	BArray< Cell_Type, Data_Type >, 38
delete_rules, 166	BArrayDense< Cell_Type, Data_Type >, 59
delete_rules_dyn, 167	swap_cols
eval rules dyn, 162	BArray< Cell_Type, Data_Type >, 39
get_counters, 163	BArrayDense< Cell Type, Data Type >, 59
get_counts, 163	swap_rows
get_counts_ptr, 163	BArray< Cell_Type, Data_Type >, 39
get_current_stats, 163	BArrayDense < Cell_Type, Data_Type >, 59
get_data, 163	target
get_rules, 164	Entries < Cell_Type >, 92
get_rules_dyn, 164	
init_support, 164	toggle_cell
M, 167	BArray< Cell_Type, Data_Type >, 39
max_num_elements, 167	BArrayDense< Cell_Type, Data_Type >, 59
N, 167	toggle_lock
print, 164	BArray< Cell_Type, Data_Type >, 39
reset_array, 164, 165	BArrayDense < Cell_Type, Data_Type >, 60
set_counters, 165	transpose
set_rules, 165	BArray< Cell_Type, Data_Type >, 39
set rules dyn, 165	BArrayDense < Cell_Type, Data_Type >, 60
Support, 160	TWO
support-meat.hpp	CHECK, 24
array_bank, 232	EXISTS, 26
BARRY_SUPPORT_MEAT_HPP, 226	typedefs.hpp
	Col_type, 237
calc_backend, 227	Counter_fun_type, 237
cfree, 232	Counts_type, 237
counters, 232	
counters_, 232	MapVec_type, 237
delete_counters, 233	Row_type, 237
delete_rules, 233	Rule_fun_type, 238
delete_rules_dyn, 233	uint, 238
else, 233	vec_equal, 238
f_, 233	vec_equal_approx, 238
for, 228	vec_inner_prod, 239
if, 228	
insert_cell, 228	ub
return, 234	PhyloRuleDynData, 141
rm_cell, 228	uint
rules, 234	typedefs.hpp, 238
rules_, 234	UKNOWN
rules_dyn, 234	EXISTS, 26
stats_bank, 234	update_annotations
SUPPORT_TEMPLATE, 227, 229–232	Geese, 111
	update_normalizing_constant
SUPPORT_TEMPLATE_ARGS, 227	model-meat.hpp, 206
SUPPORT_TYPE, 227	FF)
support_size	val
Flock, 97	Entries < Cell_Type >, 92
Geese, 111	value
Model< Array_Type, Data_Counter_Type,	Cell< Cell_Type >, 78
Data_Rule_Type, Data_Rule_Dyn_Type >,	vec_diff
126	geese-bones.hpp, 209
SUPPORT_TEMPLATE	
support-meat.hpp, 227, 229-232	vec_equal
SUPPORT_TEMPLATE_ARGS	typedefs.hpp, 238
support-meat.hpp, 227	vec_equal_approx
SUPPORT TYPE	typedefs.hpp, 238
support-meat.hpp, 227	vec_inner_prod
I- Is a	typedefs.hpp, 239

```
vecHasher < T >,\, \color{red} 168
     operator(), 168
vector_caster
     geese-bones.hpp, 209
vertex_attr
     NetworkData, 129
visited
     BArray< Cell_Type, Data_Type >, 40
     BArrayDense < Cell_Type, Data_Type >, 61
     \mathsf{Cell} \! < \mathsf{Cell} \! \_ \mathsf{Type} >, \textcolor{red}{\mathbf{78}}
     Node, 135
ZERO_CELL
     barraydense-meet.hpp, 179
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
```