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
7.1.2.2 BArray() [2/6]	30
7.1.2.3 BArray() [3/6]	30
7.1.2.4 BArray() [4/6]	31
7.1.2.5 BArray() [5/6]	31
7.1.2.6 BArray() [6/6]	31
7.1.2.7 ~BArray()	31
7.1.3 Member Function Documentation	31
7.1.3.1 clear()	31
7.1.3.2 col()	32
7.1.3.3 D() [1/2]	32
7.1.3.4 D() [2/2]	32
7.1.3.5 default_val()	32
7.1.3.6 flush_data()	32
7.1.3.7 get_cell()	32
7.1.3.8 get_col_vec() [1/2]	33
7.1.3.9 get_col_vec() [2/2]	33
7.1.3.10 get_entries()	33
7.1.3.11 get_row_vec() [1/2]	33
7.1.3.12 get_row_vec() [2/2]	33
7.1.3.13 insert_cell() [1/3]	34
7.1.3.14 insert_cell() [2/3]	34
7.1.3.15 insert_cell() [3/3]	34
7.1.3.16 is_empty()	34
7.1.3.17 ncol()	34
7.1.3.18 nnozero()	35
7.1.3.19 nrow()	35
7.1.3.20 operator()() [1/2]	35
7.1.3.21 operator()() [2/2]	35
7.1.3.22 operator*=()	35
7.1.3.23 operator+=() [1/3]	35
7.1.3.24 operator+=() [2/3]	36
7.1.3.25 operator+=() [3/3]	36
7.1.3.26 operator-=() [1/3]	36
7.1.3.27 operator-=() [2/3]	36
7.1.3.28 operator-=() [3/3]	36
7.1.3.29 operator/=()	36
7.1.3.30 operator=() [1/2]	37

7.1.3.31 operator=() [2/2]	37
7.1.3.32 operator==()	37
7.1.3.33 out_of_range()	37
7.1.3.34 print()	37
7.1.3.35 reserve()	37
7.1.3.36 resize()	38
7.1.3.37 rm_cell()	38
7.1.3.38 row()	38
7.1.3.39 set_data()	38
7.1.3.40 swap_cells()	38
7.1.3.41 swap_cols()	39
7.1.3.42 swap_rows()	39
7.1.3.43 toggle_cell()	39
7.1.3.44 toggle_lock()	39
7.1.3.45 transpose()	40
7.1.3.46 zero_col()	40
7.1.3.47 zero_row()	40
7.1.4 Friends And Related Function Documentation	40
7.1.4.1 BArrayCell < Cell_Type, Data_Type >	40
7.1.4.2 BArrayCell_const< Cell_Type, Data_Type >	40
7.1.5 Member Data Documentation	40
7.1.5.1 visited	41
7.2 BArrayCell< Cell_Type, Data_Type > Class Template Reference	41
7.2.1 Detailed Description	41
7.2.2 Constructor & Destructor Documentation	41
7.2.2.1 BArrayCell()	42
7.2.2.2 ~BArrayCell()	42
7.2.3 Member Function Documentation	42
7.2.3.1 operator Cell_Type()	42
7.2.3.2 operator*=()	42
7.2.3.3 operator+=()	42
7.2.3.4 operator-=()	43
7.2.3.5 operator/=()	43
7.2.3.6 operator=()	43
7.2.3.7 operator==()	43
7.3 BArrayCell_const< Cell_Type, Data_Type > Class Template Reference	43
7.3.1 Detailed Description	44
7.3.2 Constructor & Destructor Documentation	44
7.3.2.1 BArrayCell_const()	44
7.3.2.2 ~BArrayCell_const()	44
7.3.3 Member Function Documentation	44
7.3.3.1 operator Cell_Type()	45

7.3.3.2 operator"!=()	45
7.3.3.3 operator<()	45
7.3.3.4 operator<=()	45
7.3.3.5 operator==()	45
7.3.3.6 operator>()	46
7.3.3.7 operator>=()	46
7.4 BArrayDense < Cell_Type, Data_Type > Class Template Reference	46
7.4.1 Detailed Description	48
7.4.2 Constructor & Destructor Documentation	49
7.4.2.1 BArrayDense() [1/6]	49
7.4.2.2 BArrayDense() [2/6]	49
7.4.2.3 BArrayDense() [3/6]	49
7.4.2.4 BArrayDense() [4/6]	50
7.4.2.5 BArrayDense() [5/6]	50
7.4.2.6 BArrayDense() [6/6]	50
7.4.2.7 ~BArrayDense()	50
7.4.3 Member Function Documentation	50
7.4.3.1 clear()	51
7.4.3.2 col()	51
7.4.3.3 D() [1/2]	51
7.4.3.4 D() [2/2]	51
7.4.3.5 default_val()	51
7.4.3.6 get_cell()	52
7.4.3.7 get_col_vec() [1/2]	52
7.4.3.8 get_col_vec() [2/2]	52
7.4.3.9 get_entries()	52
7.4.3.10 get_row_vec() [1/2]	53
7.4.3.11 get_row_vec() [2/2]	53
7.4.3.12 insert_cell() [1/3]	53
7.4.3.13 insert_cell() [2/3]	53
7.4.3.14 insert_cell() [3/3]	54
7.4.3.15 is_empty()	54
7.4.3.16 ncol()	54
7.4.3.17 nnozero()	54
7.4.3.18 nrow()	54
7.4.3.19 operator()() [1/2]	55
7.4.3.20 operator()() [2/2]	55
7.4.3.21 operator*=()	55
7.4.3.22 operator+=() [1/3]	55
7.4.3.23 operator+=() [2/3]	55
7.4.3.24 operator+=() [3/3]	56
7.4.3.25 operator-=() [1/3]	56

7.4.3.26 operator-=() [2/3]	56
7.4.3.27 operator-=() [3/3]	56
7.4.3.28 operator/=()	56
7.4.3.29 operator=() [1/2]	57
7.4.3.30 operator=() [2/2]	57
7.4.3.31 operator==()	57
7.4.3.32 out_of_range()	57
7.4.3.33 print()	57
7.4.3.34 reserve()	58
7.4.3.35 resize()	58
7.4.3.36 rm_cell()	58
7.4.3.37 row()	58
7.4.3.38 set_data()	58
7.4.3.39 swap_cells()	59
7.4.3.40 swap_cols()	59
7.4.3.41 swap_rows()	59
7.4.3.42 toggle_cell()	60
7.4.3.43 toggle_lock()	60
7.4.3.44 transpose()	60
7.4.3.45 zero_col()	60
7.4.3.46 zero_row()	60
7.4.4 Friends And Related Function Documentation	61
7.4.4.1 BArrayCell< Cell_Type, Data_Type >	61
7.4.4.2 BArrayCell_const< Cell_Type, Data_Type >	61
7.4.5 Member Data Documentation	61
7.4.5.1 visited	61
$7.5 \; 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
7.9.2.2 Cell() [2/7]	75
7.9.2.3 ~Cell()	75
7.9.2.4 Cell() [3/7]	76
7.9.2.5 Cell() [4/7]	76
7.9.2.6 Cell() [5/7]	76
7.9.2.7 Cell() [6/7]	76
7.9.2.8 Cell() [7/7]	76
7.9.3 Member Function Documentation	76
7.9.3.1 add() [1/4]	77
7.9.3.2 add() [2/4]	77
7.9.3.3 add() [3/4]	77
7.9.3.4 add() [4/4]	77
7.9.3.5 operator Cell_Type()	77
7.9.3.6 operator"!=()	77
7.9.3.7 operator=() [1/2]	78
7.9.3.8 operator=() [2/2]	78
7.9.3.9 operator==()	78
7.9.4 Member Data Documentation	78
7.9.4.1 value	78
7.9.4.2 visited	78
7.10 ConstBArrayRowlter< Cell_Type, Data_Type > Class Template Reference	79
7.10.1 Detailed Description	79
7.10.2 Constructor & Destructor Documentation	79
7.10.2.1 ConstBArrayRowlter()	80
7.10.2.2 ~ConstBArrayRowlter()	80
7.10.3 Member Data Documentation	80
7.10.3.1 Array	80
7.10.3.2 current_col	80
7.10.3.3 current_row	80
7.10.3.4 iter	81
7.11 Counter< Array_Type, Data_Type > Class Template Reference	81
7.11.1 Detailed Description	82
7.11.2 Constructor & Destructor Documentation	82
7.11.2.1 Counter() [1.74]	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
7.12.2.3 Counters() [2/3]	
7.12.2.4 Counters() [3/3]	
7.12.3 Member Function Documentation	
7.12.3.1 add_counter() [1/3]	
7.12.3.2 add_counter() [2/3]	 . 88
7.12.3.3 add_counter() [3/3]	 . 88
7.12.3.4 clear()	 . 88
7.12.3.5 operator=() [1/2]	 . 88
7.12.3.6 operator=() [2/2]	 89
7.12.3.7 operator[]()	
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 set_seed()		97
7.14.3.16 support_size()		97
7.14.4 Member Data Documentation		97
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		98
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		99
7.15.3.1 add()		100
7.15.3.2 as_vector()		100
7.15.3.3 clear()		100
7.15.3.4 get_data()		100
7.15.3.5 get_data_ptr()		100
7.15.3.6 print()		101

7.15.3.7 reserve())1
7.15.3.8 size())1
7.16 Geese Class Reference)1
7.16.1 Detailed Description)3
7.16.2 Constructor & Destructor Documentation)3
7.16.2.1 Geese() [1/4])4
7.16.2.2 Geese() [2/4])4
7.16.2.3 Geese() [3/4])4
7.16.2.4 Geese() [4/4])4
7.16.2.5 ~Geese())4
7.16.3 Member Function Documentation)4
7.16.3.1 calc_reduced_sequence())5
7.16.3.2 calc_sequence())5
7.16.3.3 colnames())5
7.16.3.4 get_annotated_nodes())5
7.16.3.5 get_counters())5
7.16.3.6 get_model())5
7.16.3.7 get_probabilities())6
7.16.3.8 get_rengine())6
7.16.3.9 get_states())6
7.16.3.10 get_support())6
7.16.3.11 inherit_support())6
7.16.3.12 init())7
7.16.3.13 init_node())7
7.16.3.14 likelihood())7
7.16.3.15 likelihood_exhaust())7
7.16.3.16 nfuns())7
7.16.3.17 nleafs()	8(
7.16.3.18 nnodes()	8(
7.16.3.19 nterms())8
7.16.3.20 observed_counts())8
7.16.3.21 operator=() [1/2])8
7.16.3.22 operator=() [2/2]	8(
7.16.3.23 parse_polytomies())9
7.16.3.24 predict())9
7.16.3.25 predict_backend())9
7.16.3.26 predict_exhaust())9
7.16.3.27 predict_exhaust_backend())9
7.16.3.28 predict_sim()	0
7.16.3.29 print_observed_counts()	0
7.16.3.30 set_seed()	0
7.16.3.31 simulate()	0

7.16.3.32 support_size()	 110
7.16.3.33 update_annotations()	 111
7.16.4 Member Data Documentation	 111
7.16.4.1 delete_rengine	 111
7.16.4.2 delete_support	 111
7.16.4.3 initialized	 111
7.16.4.4 map_to_nodes	 111
7.16.4.5 nfunctions	 112
7.16.4.6 nodes	 112
7.16.4.7 reduced_sequence	 112
7.16.4.8 sequence	 112
7.17 Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_I plate Reference	 112
7.17.1 Detailed Description	
7.17.2 Constructor & Destructor Documentation	
7.17.2.1 Model() [1/3]	
7.17.2.2 Model() [2/3]	
7.17.2.3 Model() [3/3]	
7.17.2.4 ~Model()	
7.17.3 Member Function Documentation	
7.17.3.1 add_array()	
7.17.3.2 add_counter() [1/3]	
7.17.3.3 add_counter() [2/3]	
7.17.3.4 add_counter() [3/3]	 118
7.17.3.5 add_rule() [1/3]	 118
7.17.3.6 add_rule() [2/3]	 118
7.17.3.7 add_rule() [3/3]	 118
7.17.3.8 add_rule_dyn() [1/3]	 118
7.17.3.9 add_rule_dyn() [2/3]	 119
7.17.3.10 add_rule_dyn() [3/3]	 119
7.17.3.11 colnames()	 119
7.17.3.12 gen_key()	 119
7.17.3.13 get_counters()	 119
7.17.3.14 get_norm_const()	 120
7.17.3.15 get_pset()	 120
7.17.3.16 get_pset_stats()	 120
7.17.3.17 get_rengine()	 120
7.17.3.18 get_rules()	 120
7.17.3.19 get_rules_dyn()	 121
7.17.3.20 get_support()	 121
7.17.3.21 likelihood() [1/3]	 121
7.17.3.22 likelihood() [2/3]	 121

7.17.3.23 likelihood() [3/3]	 121
7.17.3.24 likelihood_total()	 122
7.17.3.25 nterms()	 122
7.17.3.26 operator=()	 122
7.17.3.27 print_stats()	 122
7.17.3.28 sample() [1/2]	 122
7.17.3.29 sample() [2/2]	 123
7.17.3.30 set_counters()	 123
7.17.3.31 set_keygen()	 123
7.17.3.32 set_rengine()	 123
7.17.3.33 set_rules()	 123
7.17.3.34 set_rules_dyn()	 124
7.17.3.35 set_seed()	 124
7.17.3.36 size()	 124
7.17.3.37 size_unique()	 124
7.17.3.38 store_psets()	 124
7.17.3.39 support_size()	 125
7.18 NetCounterData Class Reference	 125
7.18.1 Detailed Description	 125
7.18.2 Constructor & Destructor Documentation	 125
7.18.2.1 NetCounterData() [1/2]	 125
7.18.2.2 NetCounterData() [2/2]	 126
7.18.2.3 ∼NetCounterData()	 126
7.18.3 Member Data Documentation	 126
7.18.3.1 indices	 126
7.18.3.2 numbers	 126
7.19 NetworkData Class Reference	 126
7.19.1 Detailed Description	 127
7.19.2 Constructor & Destructor Documentation	 127
7.19.2.1 NetworkData() [1/3]	 127
7.19.2.2 NetworkData() [2/3]	 127
7.19.2.3 NetworkData() [3/3]	 128
7.19.2.4 ~NetworkData()	 128
7.19.3 Member Data Documentation	 128
7.19.3.1 directed	 128
7.19.3.2 vertex_attr	 129
7.20 Node Class Reference	 129
7.20.1 Detailed Description	 130
7.20.2 Constructor & Destructor Documentation	 130
7.20.2.1 Node() [1/5]	 130
7.20.2.2 Node() [2/5]	 131
7.20.2.3 Node() [3/5]	 131

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

7.22.3.10 size()	139
7.23 PhyloRuleDynData Class Reference	139
7.23.1 Detailed Description	139
7.23.2 Constructor & Destructor Documentation	139
7.23.2.1 PhyloRuleDynData()	139
$7.23.2.2 \sim$ PhyloRuleDynData()	140
7.23.3 Member Data Documentation	140
7.23.3.1 counts	140
7.23.3.2 duplication	140
7.23.3.3 lb	140
7.23.3.4 pos	140
7.23.3.5 ub	140
7.24 PowerSet< Array_Type, Data_Rule_Type > Class Template Reference	141
7.24.1 Detailed Description	142
7.24.2 Constructor & Destructor Documentation	142
7.24.2.1 PowerSet() [1/3]	142
7.24.2.2 PowerSet() [2/3]	142
7.24.2.3 PowerSet() [3/3]	143
7.24.2.4 ∼PowerSet()	143
7.24.3 Member Function Documentation	143
7.24.3.1 add_rule() [1/3]	143
7.24.3.2 add_rule() [2/3]	143
7.24.3.3 add_rule() [3/3]	143
7.24.3.4 begin()	144
7.24.3.5 calc()	144
7.24.3.6 end()	144
7.24.3.7 get_data()	144
7.24.3.8 get_data_ptr()	144
7.24.3.9 init_support()	145
7.24.3.10 operator[]()	145
7.24.3.11 reset()	145
7.24.3.12 size()	145
7.24.4 Member Data Documentation	145
7.24.4.1 coordinates_free	145
7.24.4.2 coordinates_locked	146
7.24.4.3 data	146
7.24.4.4 EmptyArray	146
7.24.4.5 M	146
7.24.4.6 N	146
7.24.4.7 rules	147
7.24.4.8 rules_deleted	147
7.25 Rule / Array Tyne, Data Tyne / Class Template Reference	1/17

7.25.1 Detailed Description	. 148
7.25.2 Constructor & Destructor Documentation	. 148
7.25.2.1 Rule() [1/2]	. 148
7.25.2.2 Rule() [2/2]	. 148
7.25.2.3 ~Rule()	. 148
7.25.3 Member Function Documentation	. 149
7.25.3.1 D()	. 149
7.25.3.2 operator()()	. 149
7.26 Rules< Array_Type, Data_Type > Class Template Reference	. 149
7.26.1 Detailed Description	. 150
7.26.2 Constructor & Destructor Documentation	. 150
7.26.2.1 Rules() [1/2]	. 150
7.26.2.2 Rules() [2/2]	. 150
7.26.2.3 ~Rules()	. 151
7.26.3 Member Function Documentation	. 151
7.26.3.1 add_rule() [1/3]	. 151
7.26.3.2 add_rule() [2/3]	. 151
7.26.3.3 add_rule() [3/3]	. 151
7.26.3.4 clear()	. 151
7.26.3.5 get_seq()	. 151
7.26.3.6 operator()()	. 152
7.26.3.7 operator=()	. 152
7.26.3.8 size()	. 153
7.27 StatsCounter< Array_Type, Data_Type > Class Template Reference	. 153
7.27.1 Detailed Description	. 153
7.27.2 Constructor & Destructor Documentation	. 154
7.27.2.1 StatsCounter() [1/2]	. 154
7.27.2.2 StatsCounter() [2/2]	. 154
7.27.2.3 ~StatsCounter()	. 154
7.27.3 Member Function Documentation	. 154
7.27.3.1 add_counter() [1/2]	. 154
7.27.3.2 add_counter() [2/2]	. 155
7.27.3.3 count_all()	. 155
7.27.3.4 count_current()	. 155
7.27.3.5 count_init()	. 155
7.27.3.6 get_counters()	. 155
7.27.3.7 reset_array()	. 155
7.27.3.8 set_counters()	. 156
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	. 159

7.	28.2.1 Support() [1/3]	. 159
7.	28.2.2 Support() [2/3]	. 159
7.	28.2.3 Support() [3/3]	. 160
7.	28.2.4 ~Support()	. 160
7.28.3 Me	ember Function Documentation	. 160
7.	28.3.1 add_counter() [1/2]	. 160
7.	28.3.2 add_counter() [2/2]	. 160
7.	28.3.3 add_rule() [1/2]	. 160
7.	28.3.4 add_rule() [2/2]	. 161
7.	28.3.5 add_rule_dyn() [1/2]	. 161
7.	28.3.6 add_rule_dyn() [2/2]	. 161
7.	28.3.7 calc()	. 161
7.	28.3.8 eval_rules_dyn()	. 162
7.	28.3.9 get_counters()	. 162
7.	28.3.10 get_counts()	. 162
7.	28.3.11 get_counts_ptr()	. 162
7.	28.3.12 get_current_stats()	. 162
7.	28.3.13 get_data()	. 163
7.	28.3.14 get_rules()	. 163
7.	28.3.15 get_rules_dyn()	. 163
7.	28.3.16 init_support()	. 163
7.	28.3.17 print()	. 163
7.	28.3.18 reset_array() [1/2]	. 164
7.	28.3.19 reset_array() [2/2]	. 164
7.	28.3.20 set_counters()	. 164
7.	28.3.21 set_rules()	. 164
7.	28.3.22 set_rules_dyn()	. 164
7.28.4 Me	ember Data Documentation	. 164
7.	28.4.1 change_stats	. 165
7.	28.4.2 coordinates_free	. 165
7.	28.4.3 coordinates_locked	. 165
7.	28.4.4 current_stats	. 165
7.	28.4.5 delete_counters	. 165
7.	28.4.6 delete_rules	. 166
7.	28.4.7 delete_rules_dyn	. 166
7.	28.4.8 M	. 166
7.	28.4.9 max_num_elements	. 166
7.	28.4.10 N	. 166
7.29 vecHasher	r< T > Struct Template Reference	. 167
7.29.1 De	tailed Description	. 167
7.29.2 Me	ember Function Documentation	. 167
7	29.2.1 operator()()	167

8 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	184
8.13.1.2 BARRY_ISFINITE	184
8.13.1.3 BARRY_MAX_NUM_ELEMENTS	184
8.13.1.4 BARRY_SAFE_EXP	184
8.13.1.5 printf_barry	184

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)1
8.21.2.4 PhyloModel)1
8.21.2.5 PhyloPowerSet)1
8.21.2.6 PhyloRule)1
8.21.2.7 PhyloRuleData)1
8.21.2.8 PhyloRuleDyn)1
8.21.2.9 PhyloRules)2
8.21.2.10 PhyloRulesDyn)2
8.21.2.11 PhyloStatsCounter)2
8.21.2.12 PhyloSupport)2
8.21.3 Function Documentation)2
8.21.3.1 get_last_name())2
8.22 include/barry/model-bones.hpp File Reference)3
8.22.1 Function Documentation)4
8.22.1.1 keygen_default())4
8.23 include/barry/model-meat.hpp File Reference)4
8.23.1 Macro Definition Documentation)5
8.23.1.1 MODEL_TEMPLATE)5
8.23.1.2 MODEL_TEMPLATE_ARGS)5
8.23.1.3 MODEL_TYPE)5
8.23.2 Function Documentation)5
8.23.2.1 likelihood_())6
8.23.2.2 MODEL_TEMPLATE() [1/2])6
8.23.2.3 MODEL_TEMPLATE() [2/2])6
8.23.2.4 update_normalizing_constant())6
8.24 include/barry/models/geese.hpp File Reference)6
8.25 include/barry/models/geese/flock-bones.hpp File Reference)7
8.26 include/barry/models/geese/flock-meet.hpp File Reference)7
8.27 include/barry/models/geese/geese-bones.hpp File Reference)8
8.27.1 Macro Definition Documentation)8
8.27.1.1 INITIALIZED)9
8.27.2 Function Documentation)9
8.27.2.1 keygen_full())9
8.27.2.2 RULE_FUNCTION())9
8.27.2.3 vec_diff())9
8.27.2.4 vector_caster())9
8.28 include/barry/models/geese/geese-meat-constructors.hpp File Reference	10
8.29 include/barry/models/geese/geese-meat-likelihood.hpp File Reference	10
8.30 include/barry/models/geese/geese-meat-likelihood_exhaust.hpp File Reference	11
8.31 include/barry/models/geese/geese-meat-predict.hpp File Reference	12
8.32 include/barry/models/geese/geese-meat-predict_exhaust.hpp File Reference	12
8.33 include/harry/models/geese/geese-meat-predict_sim_hpp_File_Reference	13

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]
8.45.2.16 SUPPORT_TEMPLATE() [9/17]
8.45.2.17 SUPPORT_TEMPLATE() [10/17]
8.45.2.18 SUPPORT_TEMPLATE() [11/17]
8.45.2.19 SUPPORT_TEMPLATE() [12/17]
8.45.2.20 SUPPORT_TEMPLATE() [13/17]
8.45.2.21 SUPPORT_TEMPLATE() [14/17]
8.45.2.22 SUPPORT_TEMPLATE() [15/17]

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

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	27
BArrayCell < Cell_Type, Data_Type >	41
BArrayCell_const< Cell_Type, Data_Type >	43
BArrayDense < Cell_Type, Data_Type >	
Baseline class for binary arrays	46
BArrayDenseCell< Cell_Type, Data_Type >	61
BArrayDenseCell_const< Cell_Type, Data_Type >	64
BArrayVector< Cell_Type, Data_Type >	
Row or column of a BArray	67
BArrayVector_const < Cell_Type, Data_Type >	71
Cell< Cell_Type >	
Entries in BArray. For now, it only has two members:	74
ConstBArrayRowlter< Cell_Type, Data_Type >	79
Counter< Array_Type, Data_Type >	
A counter function based on change statistics	81
Counters < Array_Type, Data_Type >	
Vector of counters	86
Entries < Cell_Type >	
A wrapper class to store source, target, val from a BArray object	90
Flock	
A Flock is a group of Geese	92
FreqTable < T >	
Database of statistics	98
Geese	
Annotated Phylo Model	101
Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >	
General framework for discrete exponential models. This class allows generating discrete expo-	
·	112
NetCounterData	
Data class used to store arbitrary uint or double vectors	125
NetworkData	
Data class for Networks	126
Node	
A single node for the model	129

6 Class Index

NodeData	
Data definition for the PhyloArray class	135
PhyloCounterData	136
PhyloRuleDynData	139
PowerSet< Array_Type, Data_Rule_Type >	
Powerset of a binary array	141
Rule < Array_Type, Data_Type >	
Rule for determining if a cell should be included in a sequence	147
Rules < Array_Type, Data_Type >	
Vector of objects of class Rule	149
StatsCounter< Array_Type, Data_Type >	
Count stats for a single Array	153
Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >	
Compute the support of sufficient statistics	156
vecHasher <t> 1</t>	167

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
- bool operator> (const Cell_Type &val) const
- bool operator<= (const Cell_Type &val) const
- bool operator>= (const Cell_Type &val) const

7.6.1 Detailed Description

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

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

7.6.2 Constructor & Destructor Documentation

7.6.2.1 BArrayDenseCell_const()

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

7.6.2.2 ~BArrayDenseCell_const()

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

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

7.6.3 Member Function Documentation

7.6.3.1 operator Cell_Type()

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

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

7.6.3.2 operator"!=()

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

7.6.3.3 operator<()

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

7.6.3.4 operator<=()

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

7.6.3.5 operator==()

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

7.6.3.6 operator>()

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

7.6.3.7 operator>=()

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

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

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

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

Row or column of a BArray

#include <barrayvector-bones.hpp>

Public Member Functions

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

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

7.7.1 Detailed Description

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

Row or column of a BArray

Template Parameters

Cell_Type	
Data_Type	

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

7.7.2 Constructor & Destructor Documentation

7.7.2.1 BArrayVector()

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

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

Construct a new BArrayVector object.

Parameters

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

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

7.7.2.2 ~BArrayVector()

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

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

7.7.3 Member Function Documentation

7.7.3.1 begin()

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

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

7.7.3.2 end()

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

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

7.7.3.3 is_col()

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

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

7.7.3.4 is_row()

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

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

7.7.3.5 operator std::vector< Cell_Type >()

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

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

7.7.3.6 operator*=()

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

7.7.3.7 operator+=()

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

7.7.3.8 operator-=()

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

7.7.3.9 operator/=()

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

7.7.3.10 operator=()

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

7.7.3.11 operator==()

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

7.7.3.12 size()

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

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

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

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

7.8 BArrayVector_const< Cell_Type, Data_Type > Class Template Reference

#include <barrayvector-bones.hpp>

Public Member Functions

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

7.8.1 Detailed Description

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

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

7.8.2 Constructor & Destructor Documentation

7.8.2.1 BArrayVector_const()

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

7.8.2.2 ~BArrayVector_const()

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

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

7.8.3 Member Function Documentation

7.8.3.1 begin()

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

7.8.3.2 end()

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

7.8.3.3 is col()

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

7.8.3.4 is_row()

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

7.8.3.5 operator std::vector< Cell_Type >()

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

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

7.8.3.6 operator"!=()

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

7.8.3.7 operator<()

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

7.8.3.8 operator<=()

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

7.8.3.9 operator==()

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

7.8.3.10 operator>()

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

7.8.3.11 operator>=()

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

7.8.3.12 size()

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

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

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

7.9 Cell < Cell Type > Class Template Reference

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

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

Public Member Functions

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

Public Attributes

- Cell_Type value
- bool visited

7.9.1 Detailed Description

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

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

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

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

7.9.2 Constructor & Destructor Documentation

7.9.2.1 Cell() [1/7]

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

7.9.2.2 Cell() [2/7]

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

7.9.2.3 ∼CeII()

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

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

7.9.2.4 Cell() [3/7]

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

7.9.2.5 Cell() [4/7]

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

7.9.2.6 Cell() [5/7]

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

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

7.9.2.7 Cell() [6/7]

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

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

7.9.2.8 Cell() [7/7]

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

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

7.9.3 Member Function Documentation

7.9.3.1 add() [1/4]

7.9.3.2 add() [2/4]

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

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

7.9.3.3 add() [3/4]

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

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

7.9.3.4 add() [4/4]

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

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

7.9.3.5 operator Cell_Type()

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

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

7.9.3.6 operator"!=()

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

7.9.3.7 operator=() [1/2]

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

7.9.3.8 operator=() [2/2]

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

7.9.3.9 operator==()

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

7.9.4 Member Data Documentation

7.9.4.1 value

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

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

7.9.4.2 visited

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

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

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

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

7.10 ConstBArrayRowlter< Cell_Type, Data_Type > Class Template Reference

#include <barray-iterator.hpp>

Collaboration diagram for ConstBArrayRowlter< Cell_Type, Data_Type >:



Public Member Functions

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

Public Attributes

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

7.10.1 Detailed Description

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

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

7.10.2 Constructor & Destructor Documentation

7.10.2.1 ConstBArrayRowlter()

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

7.10.2.2 ~ConstBArrayRowlter()

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

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

7.10.3 Member Data Documentation

7.10.3.1 Array

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

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

7.10.3.2 current_col

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

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

7.10.3.3 current_row

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

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

7.10.3.4 iter

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

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

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

include/barry/barray-iterator.hpp

7.11 Counter< Array_Type, Data_Type > Class Template Reference

A counter function based on change statistics.

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

Public Member Functions

- ∼Counter ()
- double count (Array_Type &Array, uint i, uint j)
- double init (Array_Type &Array, uint i, uint j)

Creator passing a counter and an initializer

Parameters

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

- Counter ()
- Counter (Counter_fun_type< Array_Type, Data_Type > count_fun_, Counter_fun_type< Array_Type, Data_Type > init_fun_=nullptr, Data_Type *data_=nullptr, bool delete_data_=false, std::string name_
 ="", std::string desc ="")
- Counter (const Counter < Array_Type, Data_Type > &counter_)
 Copy constructor.
- Counter (Counter < Array_Type, Data_Type > &&counter_) noexcept
 Move constructor.
- Counter< Array_Type, Data_Type > operator= (const Counter< Array_Type, Data_Type > &counter_)
 Copy assignment.
- Counter< Array_Type, Data_Type > & operator= (Counter< Array_Type, Data_Type > &&counter_←) noexcept

Move assignment.

Public Attributes

```
    Counter_fun_type< Array_Type, Data_Type > count_fun
    Counter_fun_type< Array_Type, Data_Type > init_fun
    Data_Type * data = nullptr
    bool delete_data = false
    std::string name = ""
    std::string desc = ""
```

7.11.1 Detailed Description

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

A counter function based on change statistics.

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

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

7.11.2 Constructor & Destructor Documentation

7.11.2.1 Counter() [1/4]

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

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

7.11.2.2 Counter() [2/4]

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

7.11.2.3 Counter() [3/4]

Copy constructor.

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

7.11.2.4 Counter() [4/4]

Move constructor.

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

7.11.2.5 ~Counter()

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

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

7.11.3 Member Function Documentation

7.11.3.1 count()

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

7.11.3.2 init()

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

7.11.3.3 operator=() [1/2]

Copy assignment.

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

7.11.3.4 operator=() [2/2]

Move assignment.

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

7.11.4 Member Data Documentation

7.11.4.1 count_fun

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

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

7.11.4.2 data

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

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

7.11.4.3 delete_data

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

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

7.11.4.4 desc

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

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

7.11.4.5 init_fun

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

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

7.11.4.6 name

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

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

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

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

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

Vector of counters.

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

Public Member Functions

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

Copy constructor.

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

Move constructor.

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

Move assignment constructor.

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

Returns a pointer to a particular counter.

• std::size_t size () const noexcept

Number of counters in the set.

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

7.12.1 Detailed Description

Vector of counters.

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

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

7.12.2 Constructor & Destructor Documentation

7.12.2.1 Counters() [1/3]

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

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

7.12.2.2 ∼Counters()

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

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

7.12.2.3 Counters() [2/3]

Copy constructor.

Parameters



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

7.12.2.4 Counters() [3/3]

Move constructor.

Parameters



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

7.12.3 Member Function Documentation

7.12.3.1 add counter() [1/3]

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

7.12.3.2 add_counter() [2/3]

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

7.12.3.3 add_counter() [3/3]

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

7.12.3.4 clear()

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

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

7.12.3.5 operator=() [1/2]

Copy assignment constructor.

Parameters

counter←	
_	

Returns

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

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

7.12.3.6 operator=() [2/2]

Move assignment constructor.

Parameters



Returns

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

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

7.12.3.7 operator[]()

Returns a pointer to a particular counter.

Parameters

```
idx Id of the counter
```

Returns

Counter<Array_Type,Data_Type>*

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

7.12.3.8 size()

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

Number of counters in the set.

Returns

uint

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

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

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

7.13 Entries < Cell_Type > Class Template Reference

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

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

Public Member Functions

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

Public Attributes

- std::vector< uint > source
- std::vector< uint > target
- std::vector< Cell Type > val

7.13.1 Detailed Description

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

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

Template Parameters

```
Cell_Type Any type
```

Definition at line 59 of file typedefs.hpp.

7.13.2 Constructor & Destructor Documentation

7.13.2.1 Entries() [1/2]

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

Definition at line 65 of file typedefs.hpp.

7.13.2.2 Entries() [2/2]

Definition at line 66 of file typedefs.hpp.

7.13.2.3 ∼Entries()

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

Definition at line 73 of file typedefs.hpp.

7.13.3 Member Function Documentation

7.13.3.1 resize()

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

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

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.

Parameters

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

Returns

Geese *

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

7.14.3.14 parse_polytomies()

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

7.14.3.15 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.

7.14.3.16 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.

7.15.3.6 print()

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

Definition at line 102 of file statsdb.hpp.

7.15.3.7 reserve()

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 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 ${\tt N}.$
parent	Id of the parent gene. Also of length ${\tt N}$

- 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 < std::string > colnames () const

Names of the terms in the model.

• unsigned int parse_polytomies (bool verb=true) const noexcept

Check polytomies and return the largest.

Geese prediction

Calculate the conditional probability

Parameters

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

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

Returns

std::vector< double > Returns the posterior probability

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

 annotated=false, unsigned int nsims=10000u)

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

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

Returns

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

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

Public Attributes

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

7.16.1 Detailed Description

Annotated Phylo Model.

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

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

7.16.2 Constructor & Destructor Documentation

7.16.2.1 Geese() [1/4]

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

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

7.16.2.2 Geese() [2/4]

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

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

7.16.2.3 Geese() [3/4]

Definition at line 180 of file geese-meat-constructors.hpp.

7.16.2.4 Geese() [4/4]

Definition at line 249 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 Geese Class Reference 105

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 367 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 527 of file geese-meat.hpp.

7.16.3.5 get_counters()

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

Definition at line 511 of file geese-meat.hpp.

7.16.3.6 get_model()

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

Definition at line 515 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 507 of file geese-meat.hpp.

7.16.3.9 get_states()

```
\verb|std::vector| < \verb|std::vector| < \verb|std::vector| > > Geese::get_states () const [inline]|
```

Powerset of a gene's possible states.

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

Returns

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

Definition at line 523 of file geese-meat.hpp.

7.16.3.10 get_support()

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

Definition at line 519 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()

```
void Geese::init (
          bool verb = true ) [inline]
```

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

7.16.3.13 init_node()

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

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 nfuns()

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

Number of functions analyzed.

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

7.16.3.17 nleafs()

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

Number of leaf.

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

7.16.3.18 nnodes()

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

Number of nodes (interior + leaf)

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

7.16.3.19 nterms()

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

Number of terms included.

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

7.16.3.20 observed_counts()

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

Definition at line 397 of file geese-meat.hpp.

7.16.3.21 operator=() [1/2]

7.16.3.22 operator=() [2/2]

7.16 Geese Class Reference 109

7.16.3.23 parse_polytomies()

Check polytomies and return the largest.

Definition at line 373 of file geese-meat.hpp.

7.16.3.24 predict()

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

7.16.3.25 predict backend()

< True if the array belongs to the set

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

7.16.3.26 predict_exhaust()

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

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

7.16.3.27 predict_exhaust_backend()

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

7.16.3.28 predict_sim()

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

7.16.3.29 print_observed_counts()

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

Definition at line 445 of file geese-meat.hpp.

7.16.3.30 set_seed()

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

7.16.3.31 simulate()

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

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

7.16.3.32 support_size()

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

Number of unique sets of sufficient stats.

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

7.16 Geese Class Reference 111

7.16.3.33 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 113 of file geese-bones.hpp.

7.16.4.2 delete_support

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

Definition at line 114 of file geese-bones.hpp.

7.16.4.3 initialized

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

Definition at line 112 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 105 of file geese-bones.hpp.

7.16.4.5 nfunctions

unsigned int Geese::nfunctions

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

7.16.4.6 nodes

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

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

7.16.4.7 reduced_sequence

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

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

7.16.4.8 sequence

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

Definition at line 108 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
- Array_Type sample (const Array_Type &Array_, const std::vector< double > ¶ms={})
- Array Type sample (const uint &i, const std::vector< double > ¶ms)
- const std::mt19937 * get rengine () const
- Counters < Array Type, Data Counter Type > * get counters ()
- Rules
 Array_Type, Data_Rule_Type > * get_rules ()
- Rules< Array_Type, Data_Rule_Dyn_Type > * get_rules_dyn ()
- Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > * get_support ()

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

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

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

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

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

- void add_rule (Rule < Array_Type, Data_Rule_Type > &rule)
- void add rule (Rule < Array Type, Data Rule Type > *rule)
- void add_rule (Rule_fun_type< Array_Type, Data_Rule_Type > count_fun_, Data_Rule_Type *data_←
 =nullptr, bool delete_data_=false)
- void set_rules (Rules< Array_Type, Data_Rule_Type > *rules_)
- void add_rule_dyn (Rule< Array_Type, Data_Rule_Dyn_Type > &rule)
- void add_rule_dyn (Rule< Array_Type, Data_Rule_Dyn_Type > *rule)
- void 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 > ¶ms, const uint &i, bool as_log=false)
- double likelihood (const std::vector< double > ¶ms, const std::vector< double > &target_, const uint &i, bool as log=false)
- double likelihood_total (const std::vector< double > ¶ms, bool as_log=false)

Extract elements by index

Parameters

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

- double get norm const (const std::vector< double > ¶ms, const uint &i, bool as log=false)
- const std::vector< Array Type > * get pset (const uint &i)
- const std::vector< std::vector< double > > * get_pset_stats (const uint &i)

Size of the model

Number of different supports included in the model

This will return the size of stats.

Returns

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

- unsigned int size () const noexcept
- unsigned int size_unique () const noexcept
- unsigned int nterms () const noexcept
- unsigned int support_size () const noexcept
- std::vector< std::string > colnames () const

7.17.1 Detailed Description

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

class Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >

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

$$\frac{\exp\left(\theta^{\dagger}c(A)\right)}{\sum_{A'\in\mathcal{A}}\exp\left(\theta^{\dagger}c(A')\right)}$$

This implementation aims to reduce the number of times that the support needs to be computed. Models included here use more than a single array, and thus allow the function to recycle support sets as needed. For example, if we are looking at directed graphs all of the same size and without vertex level features, i.e. a model that only counts edges, triangles, etc. then the support needs to be fully computed only once.

Template Parameters

Array_Type	Class of BArray object.
Data_Counter_Type	Any type.
Data_Rule_Type	Any type.

Definition at line 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()

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

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_stats()

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

7.17.3.28 sample() [1/2]

7.17.3.29 sample() [2/2]

7.17.3.30 set_counters()

7.17.3.31 set_keygen()

7.17.3.32 set_rengine()

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

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

7.17.3.33 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_ )
```

7.17.3.34 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_ )
```

7.17.3.35 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.

7.17.3.36 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]
```

7.17.3.37 size_unique()

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

7.17.3.38 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]
```

7.17.3.39 support_size()

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

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

• include/barry/model-bones.hpp

7.18 NetCounterData Class Reference

Data class used to store arbitrary uint or double vectors.

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

Public Member Functions

- NetCounterData ()
- NetCounterData (const std::vector< uint > indices_, const std::vector< double > numbers_)
- ∼NetCounterData ()

Public Attributes

- std::vector< uint > indices
- std::vector< double > numbers

7.18.1 Detailed Description

Data class used to store arbitrary uint or double vectors.

Definition at line 61 of file network.hpp.

7.18.2 Constructor & Destructor Documentation

7.18.2.1 NetCounterData() [1/2]

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

Definition at line 67 of file network.hpp.

7.18.2.2 NetCounterData() [2/2]

Definition at line 68 of file network.hpp.

7.18.2.3 ∼NetCounterData()

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

Definition at line 73 of file network.hpp.

7.18.3 Member Data Documentation

7.18.3.1 indices

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

Definition at line 64 of file network.hpp.

7.18.3.2 numbers

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

Definition at line 65 of file network.hpp.

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

• include/barry/counters/network.hpp

7.19 NetworkData Class Reference

Data class for Networks.

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

Public Member Functions

- · NetworkData ()
- NetworkData (std::vector< double > vertex_attr_, bool directed_=true)

Constructor using a single attribute.

NetworkData (std::vector< std::vector< double >> vertex_attr_, bool directed_=true)

Constructor using multiple attributes.

∼NetworkData ()

Public Attributes

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

7.19.1 Detailed Description

Data class for Networks.

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

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

Definition at line 24 of file network.hpp.

7.19.2 Constructor & Destructor Documentation

7.19.2.1 NetworkData() [1/3]

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

Definition at line 30 of file network.hpp.

7.19.2.2 NetworkData() [2/3]

Constructor using a single attribute.

Parameters

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

Definition at line 38 of file network.hpp.

7.19.2.3 NetworkData() [3/3]

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

Constructor using multiple attributes.

Parameters

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

Definition at line 50 of file network.hpp.

7.19.2.4 ~NetworkData()

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

Definition at line 56 of file network.hpp.

7.19.3 Member Data Documentation

7.19.3.1 directed

bool NetworkData::directed = true

Definition at line 27 of file network.hpp.

7.20 Node Class Reference 129

7.19.3.2 vertex_attr

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

Definition at line 28 of file network.hpp.

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

• include/barry/counters/network.hpp

7.20 Node Class Reference

A single node for the model.

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

Collaboration diagram for Node:



Public Member Functions

- ∼Node ()
- int get_parent () const
- unsigned int noffspring () const noexcept
- bool is_leaf () const noexcept

Construct a new Node object

- Node ()
- Node (unsigned int id_, unsigned int ord_, bool duplication_)
- Node (unsigned int id_, unsigned int ord_, std::vector< unsigned int > annotations_, bool duplication_)
- Node (Node &&x) noexcept
- Node (const Node &x)

Public Attributes

· unsigned int id

Id of the node (as specified in the input)

· unsigned int ord

Order in which the node was created.

- · phylocounters::PhyloArray array
- std::vector< unsigned int > annotations

Observed annotations (only defined for Geese)

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

Arrays given all possible states.

Node * parent = nullptr

Parent node.

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

Offspring nodes.

• std::vector< unsigned int > narray = {}

ID of the array in the model.

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

Induced subtree probabilities.

std::vector< double > probability

The probability of observing each state.

7.20.1 Detailed Description

A single node for the model.

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

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

7.20.2 Constructor & Destructor Documentation

7.20.2.1 Node() [1/5]

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

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

7.20 Node Class Reference 131

7.20.2.2 Node() [2/5]

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

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

7.20.2.3 Node() [3/5]

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

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

7.20.2.4 Node() [4/5]

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

7.20.2.5 Node() [5/5]

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

7.20.2.6 ∼Node()

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

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

7.20.3 Member Function Documentation

7.20.3.1 get_parent()

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

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

7.20.3.2 is_leaf()

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

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

7.20.3.3 noffspring()

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

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

7.20.4 Member Data Documentation

7.20.4.1 annotations

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

Observed annotations (only defined for Geese)

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

7.20.4.2 array

phylocounters::PhyloArray Node::array

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

7.20 Node Class Reference 133

7.20.4.3 arrays

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

Arrays given all possible states.

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

7.20.4.4 duplication

```
bool Node::duplication
```

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

7.20.4.5 id

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

Id of the node (as specified in the input)

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

7.20.4.6 narray

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

ID of the array in the model.

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

7.20.4.7 offspring

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

Offspring nodes.

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

7.20.4.8 ord

```
unsigned int Node::ord
```

Order in which the node was created.

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

7.20.4.9 parent

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

Parent node.

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

7.20.4.10 probability

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

The probability of observing each state.

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

7.20.4.11 subtree_prob

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

Induced subtree probabilities.

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

7.20.4.12 visited

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

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

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

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

7.21 NodeData Class Reference

Data definition for the PhyloArray class.

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

Public Member Functions

NodeData (const std::vector< double > &blengths_, const std::vector< bool > &states_, bool duplication
 —=true)

Public Attributes

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

7.21.1 Detailed Description

Data definition for the PhyloArray class.

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

This holds basic information about a given node.

Definition at line 23 of file phylo.hpp.

7.21.2 Constructor & Destructor Documentation

7.21.2.1 NodeData()

Definition at line 43 of file phylo.hpp.

7.21.3 Member Data Documentation

7.21.3.1 blengths

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

Branch length.

Definition at line 29 of file phylo.hpp.

7.21.3.2 duplication

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

Definition at line 39 of file phylo.hpp.

7.21.3.3 states

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

State of the parent node.

Definition at line 34 of file phylo.hpp.

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

• include/barry/counters/phylo.hpp

7.22 PhyloCounterData Class Reference

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

Public Member Functions

- PhyloCounterData (std::vector< uint > data_, std::vector< double > *counters_=nullptr)
- uint at (uint d)
- uint operator() (uint d)
- void reserve (uint x)
- void push_back (uint x)
- void shrink_to_fit ()
- uint size ()
- std::vector< uint >::iterator begin ()
- std::vector< uint >::iterator end ()
- bool empty ()
- std::vector< double > * get_counters ()

7.22.1 Detailed Description

Definition at line 54 of file phylo.hpp.

7.22.2 Constructor & Destructor Documentation

7.22.2.1 PhyloCounterData()

Definition at line 60 of file phylo.hpp.

7.22.3 Member Function Documentation

7.22.3.1 at()

Definition at line 65 of file phylo.hpp.

7.22.3.2 begin()

```
\verb|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:

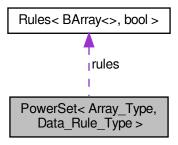
• include/barry/counters/phylo.hpp

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.

- ∼StatsCounter ()
- void reset_array (const Array_Type *Array_)

Changes the reference array for the counting.

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

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

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

7.27.1 Detailed Description

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

Count stats for a single Array.

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

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

7.27.2 Constructor & Destructor Documentation

7.27.2.1 StatsCounter() [1/2]

Creator of a StatsCounter

Parameters

Array←	A const pointer to a BArray.	
_		

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

7.27.2.2 StatsCounter() [2/2]

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

Can be created without setting the array.

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

7.27.2.3 ~StatsCounter()

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

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

7.27.3 Member Function Documentation

7.27.3.1 add_counter() [1/2]

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

7.27.3.2 add_counter() [2/2]

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

7.27.3.3 count all()

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

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

7.27.3.4 count_current()

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

7.27.3.5 count init()

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

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

7.27.3.6 get_counters()

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

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

7.27.3.7 reset_array()

Changes the reference array for the counting.

Parameters

Array⊷	A pointer to an array of class Array_Type.

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

7.27.3.8 set_counters()

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

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

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

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

Compute the support of sufficient statistics.

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

Public Member Functions

Support (const Array_Type &Array_)

Constructor passing a reference Array.

Support (uint N_, uint M_)

Constructor specifying the dimensions of the array (empty).

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

Computes the entire support.

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

List current statistics.

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

Vector of couter functions.

• Rules< Array_Type, Data_Rule_Type > * get_rules ()

Vector of static rules (cells to iterate).

Rules< Array_Type, Data_Rule_Dyn_Type > * get_rules_dyn ()

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

Resets the support calculator

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

158 Class Documentation

Parameters

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

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

Manage counters

Parameters

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

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

Manage rules

Parameters

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

- void add_rule (Rule < Array_Type, Data_Rule_Type > *f_)
- void add_rule (Rule < Array_Type, Data_Rule_Type > f_)
- void set_rules (Rules< Array_Type, Data_Rule_Type > *rules_)
- void add_rule_dyn (Rule < Array_Type, Data_Rule_Dyn_Type > *f_)
- void add_rule_dyn (Rule < Array_Type, Data_Rule_Dyn_Type > f_)
- void set_rules_dyn (Rules < Array_Type, Data_Rule_Dyn_Type > *rules_)
- bool eval_rules_dyn (const std::vector< double > &counts, const uint &i, const uint &j)

Public Attributes

- uint N
- uint M
- bool delete_counters = true
- bool delete_rules = true
- bool delete_rules_dyn = true
- uint max_num_elements = BARRY_MAX_NUM_ELEMENTS
- $\bullet \ \, \mathsf{std} : \! \mathsf{vector} \! < \mathsf{double} > \! \mathsf{current_stats}$
- std::vector< std::pair< uint, uint >> coordinates_free
- std::vector< std::pair< uint, uint >> coordinates_locked
- std::vector< std::vector< double >> change stats

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.

160 Class Documentation

7.28.2.3 Support() [3/3]

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

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

7.28.2.4 ~Support()

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

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

7.28.3 Member Function Documentation

7.28.3.1 add counter() [1/2]

7.28.3.2 add_counter() [2/2]

7.28.3.3 add_rule() [1/2]

7.28.3.4 add_rule() [2/2]

7.28.3.5 add_rule_dyn() [1/2]

7.28.3.6 add_rule_dyn() [2/2]

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

7.28.3.7 calc()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::calc (
    std::vector< Array_Type > * array_bank = nullptr,
    std::vector< std::vector< double > > * stats_bank = nullptr,
    unsigned int max_num_elements_ = 0u )
```

Computes the entire support.

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

Parameters

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

162 Class Documentation

7.28.3.8 eval_rules_dyn()

7.28.3.9 get_counters()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Counters<Array_Type, Data_Counter_Type>* Support< Array_Type, Data_Counter_Type, Data_Rule_←
Type, Data_Rule_Dyn_Type >::get_counters ()
```

Vector of couter functions.

7.28.3.10 get_counts()

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

7.28.3.11 get_counts_ptr()

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

7.28.3.12 get_current_stats()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< double >* Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_←
Dyn_Type >::get_current_stats ()
```

List current statistics.

7.28.3.13 get_data()

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

7.28.3.14 get_rules()

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

Vector of static rules (cells to iterate).

7.28.3.15 get_rules_dyn()

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

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

7.28.3.16 init support()

7.28.3.17 print()

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

164 Class Documentation

7.28.3.18 reset_array() [1/2]

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

7.28.3.19 reset_array() [2/2]

7.28.3.20 set_counters()

7.28.3.21 set_rules()

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

7.28.3.22 set_rules_dyn()

7.28.4 Member Data Documentation

7.28.4.1 change_stats

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

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

7.28.4.2 coordinates free

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

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

7.28.4.3 coordinates locked

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

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

7.28.4.4 current_stats

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

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

7.28.4.5 delete_counters

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

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

166 Class Documentation

7.28.4.6 delete_rules

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

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

7.28.4.7 delete_rules_dyn

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

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

7.28.4.8 M

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

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

7.28.4.9 max_num_elements

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

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

7.28.4.10 N

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

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

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

• include/barry/support-bones.hpp

7.29 vecHasher< T > Struct Template Reference

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

Public Member Functions

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

7.29.1 Detailed Description

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

Definition at line 86 of file typedefs.hpp.

7.29.2 Member Function Documentation

7.29.2.1 operator()()

Definition at line 87 of file typedefs.hpp.

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

• include/barry/typedefs.hpp

168 Class Documentation

Chapter 8

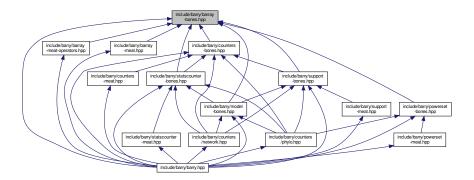
File Documentation

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

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



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



Classes

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

Macros

• #define BARRAY_BONES_HPP 1

8.1.1 Macro Definition Documentation

8.1.1.1 BARRAY_BONES_HPP

#define BARRAY_BONES_HPP 1

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

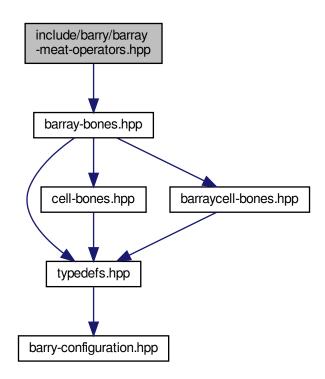
8.2 include/barry/barray-iterator.hpp File Reference

Classes

class ConstBArrayRowIter< Cell_Type, Data_Type >

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

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



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



Macros

- #define BARRY_BARRAY_MEAT_OPERATORS_HPP 1
- #define ROW(a) this->el_ij[a]
- #define COL(a) this->el_ji[a]

Functions

template<typename Cell_Type , typename Data_Type >
 void checkdim_ (const BArray< Cell_Type, Data_Type > &lhs, const BArray< Cell_Type, Data_Type > &rhs)

8.3.1 Macro Definition Documentation

8.3.1.1 BARRY_BARRAY_MEAT_OPERATORS_HPP

```
#define BARRY_BARRAY_MEAT_OPERATORS_HPP 1
```

Definition at line 5 of file barray-meat-operators.hpp.

8.3.1.2 COL

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

8.3.1.3 ROW

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

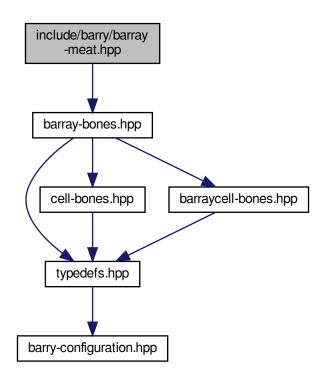
8.3.2 Function Documentation

8.3.2.1 checkdim_()

Definition at line 11 of file barray-meat-operators.hpp.

8.4 include/barry/barray-meat.hpp File Reference

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



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



Macros

- #define ROW(a) this->el_ij[a]
- #define COL(a) this->el_ji[a]

8.4.1 Macro Definition Documentation

8.4.1.1 COL

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

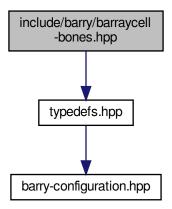
8.4.1.2 ROW

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

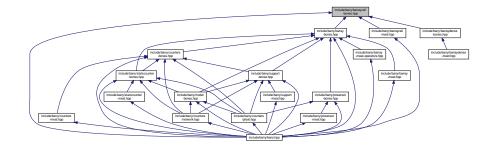
Definition at line 7 of file barray-meat.hpp.

8.5 include/barry/barraycell-bones.hpp File Reference

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



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

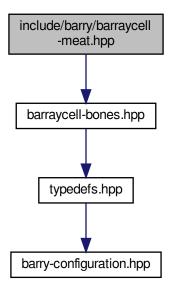


Classes

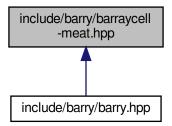
- class BArrayCell
 Cell_Type, Data_Type
- class BArrayCell_const< Cell_Type, Data_Type >

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

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



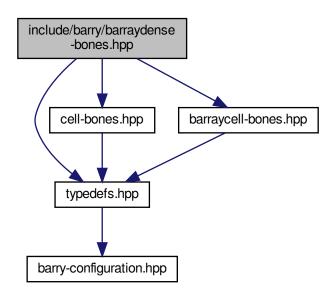
This graph shows which files directly or indirectly include this file:



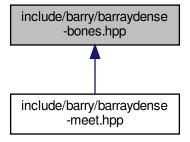
8.7 include/barry/barraydense-bones.hpp File Reference

```
#include "typedefs.hpp"
#include "cell-bones.hpp"
#include "barraycell-bones.hpp"
```

Include dependency graph for barraydense-bones.hpp:



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



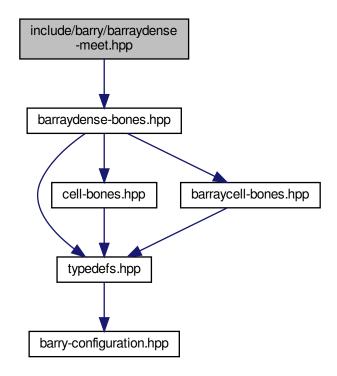
Classes

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

8.8 include/barry/barraydense-meet.hpp File Reference

#include "barraydense-bones.hpp"

Include dependency graph for barraydense-meet.hpp:



Macros

- #define BARRY_BARRAYDENSE_MEAT_HPP
- #define ROW(a) this->el_ij[a]
- #define COL(a) this->el_ji[a]
- #define POS(a, b) (b)*N + (a)
- #define ZERO_CELL Cell< Cell_Type >(static_cast< Cell_Type >(0.0))

8.8.1 Macro Definition Documentation

8.8.1.1 BARRY_BARRAYDENSE_MEAT_HPP

#define BARRY_BARRAYDENSE_MEAT_HPP

Definition at line 5 of file barraydense-meet.hpp.

8.8.1.2 COL

```
#define COL( a \ ) \ \  \  this->el_ji[a]
```

Definition at line 8 of file barraydense-meet.hpp.

8.8.1.3 POS

```
#define POS(  a, \\ b ) \ (b)*N + (a)
```

Definition at line 9 of file barraydense-meet.hpp.

8.8.1.4 ROW

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

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

8.8.1.5 ZERO_CELL

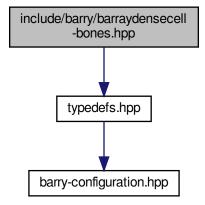
```
#define ZERO_CELL Cell Cell_Type >(static_cast< Cell_Type >(0.0))
```

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

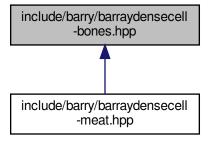
8.9 include/barry/barraydensecell-bones.hpp File Reference

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

Include dependency graph for barraydensecell-bones.hpp:



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

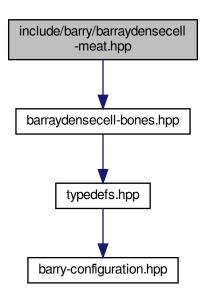


Classes

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

8.10 include/barry/barraydensecell-meat.hpp File Reference

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



Macros

- #define BARRY_BARRAYDENSECELL_MEAT_HPP 1
- #define POS(a, b) (a) + (b) * Array->N

8.10.1 Macro Definition Documentation

8.10.1.1 BARRY_BARRAYDENSECELL_MEAT_HPP

```
#define BARRY_BARRAYDENSECELL_MEAT_HPP 1
```

Definition at line 4 of file barraydensecell-meat.hpp.

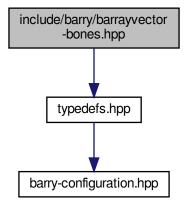
8.10.1.2 POS

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

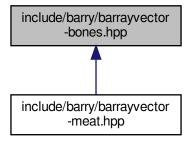
Definition at line 6 of file barraydensecell-meat.hpp.

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:

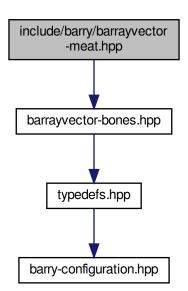


Classes

- 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

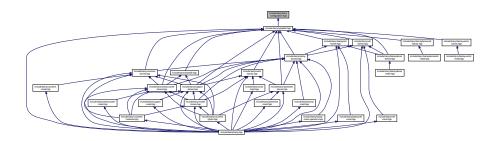
8.12.1.1 BARRY BARRAYVECTOR MEAT HPP

#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

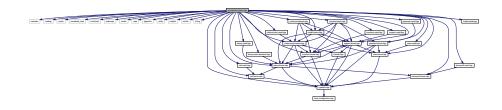
8.13.2.1 Map

```
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

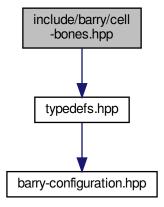
Definition at line 80 of file barry.hpp.

Definition at line 83 of file barry.hpp.

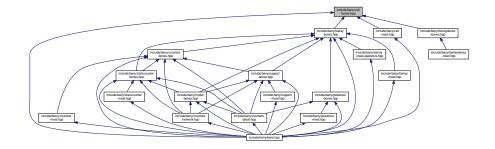
8.14.1.6 RULE_LAMBDA

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:

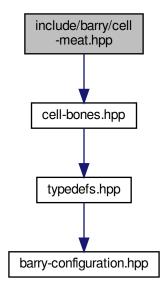


Classes

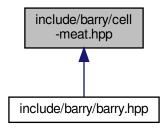
class Cell
 Cell_Type >
 Entries in BArray. For now, it only has two members:

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

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



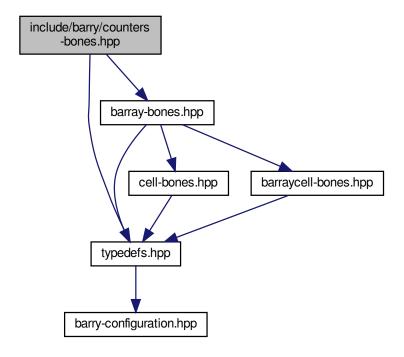
This graph shows which files directly or indirectly include this file:



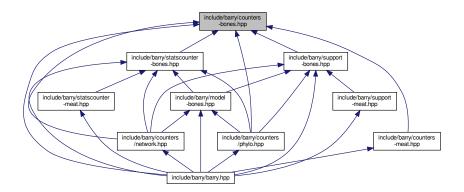
8.17 include/barry/col-bones.hpp File Reference

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:



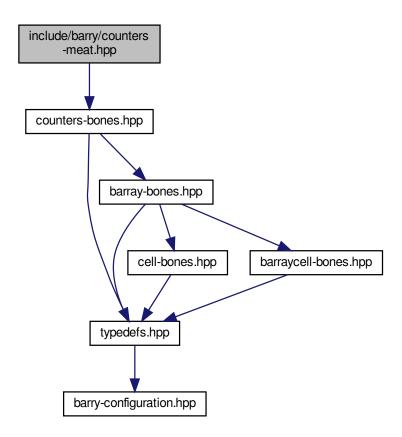
Classes

- 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:



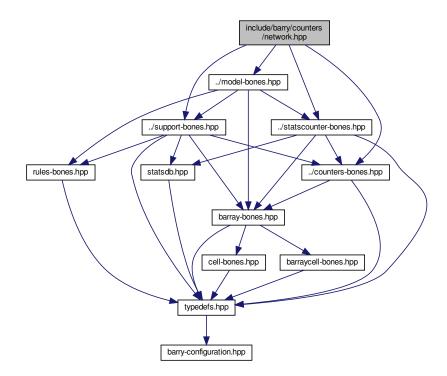
This graph shows which files directly or indirectly include this file:



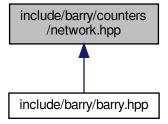
8.20 include/barry/counters/network.hpp File Reference

```
#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

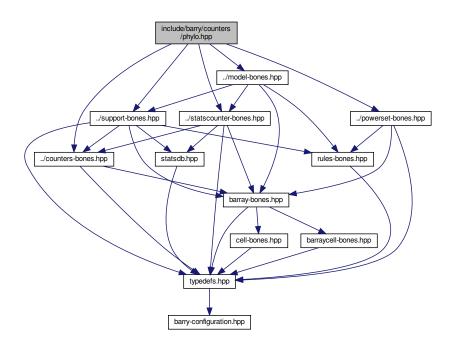
8.20.3.1 rules_zerodiag()

Number of edges.

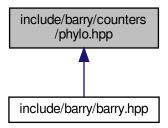
Definition at line 803 of file network.hpp.

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

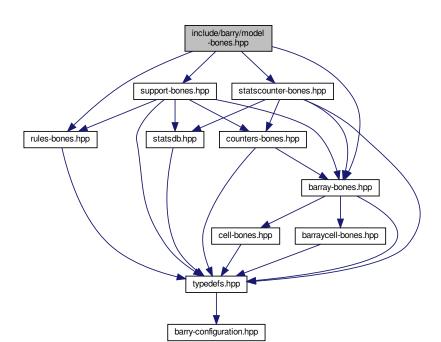
8.21.3.1 get_last_name()

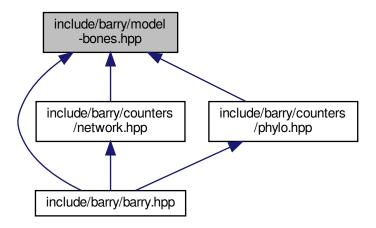
```
std::string get_last_name (
          bool d ) [inline]
```

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

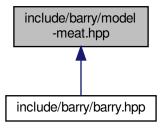
8.22.1.1 keygen_default()

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 > ¶ms, const Counts_type &support)
- double likelihood_ (const std::vector< double > &target_stats, const std::vector< double > ¶ms, const double normalizing_constant, bool log_=false)
- MODEL_TEMPLATE (, Model)()
- MODEL_TEMPLATE (, Model)(const MODEL_TYPE() &Model_)

8.23.1 Macro Definition Documentation

8.23.1.1 MODEL_TEMPLATE

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

8.23.1.2 MODEL_TEMPLATE_ARGS

```
#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:

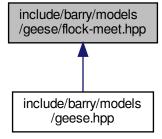


Classes

· 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

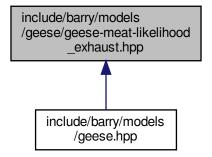
#include "geese-bones.hpp"
Include dependency graph for geese-meat-likelihood.hpp:



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:

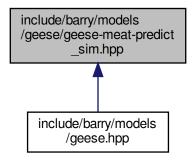


8.32 include/barry/models/geese/geese-meat-predict_exhaust.hpp File Reference



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

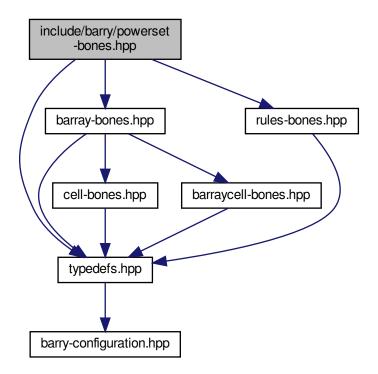
• class Node

A single node for the model.

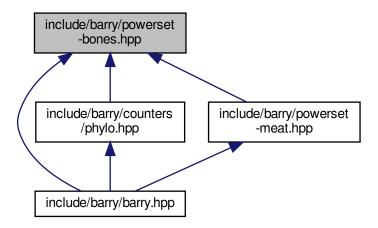
8.37 include/barry/powerset-bones.hpp File Reference

```
#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:

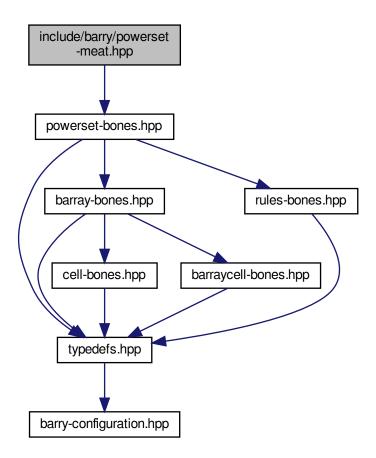


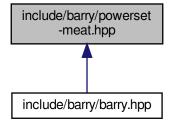
Classes

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

#include "powerset-bones.hpp"

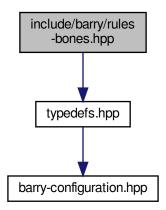
Include dependency graph for powerset-meat.hpp:



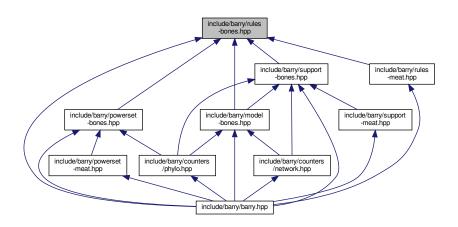


8.39 include/barry/rules-bones.hpp File Reference

#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)

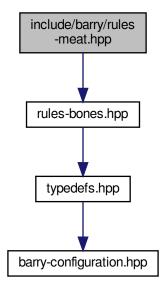
8.39.1 Function Documentation

8.39.1.1 rule_fun_default()

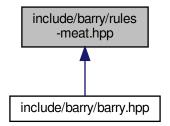
Definition at line 10 of file rules-bones.hpp.

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

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



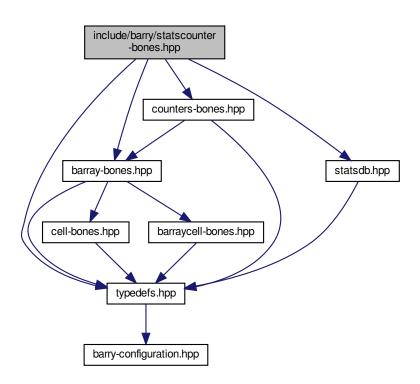
This graph shows which files directly or indirectly include this file:



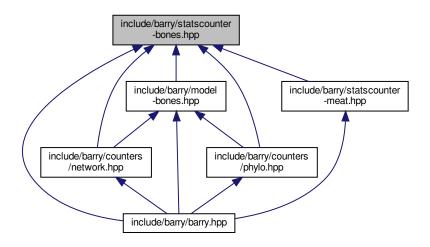
8.41 include/barry/statscounter-bones.hpp File Reference

```
#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:



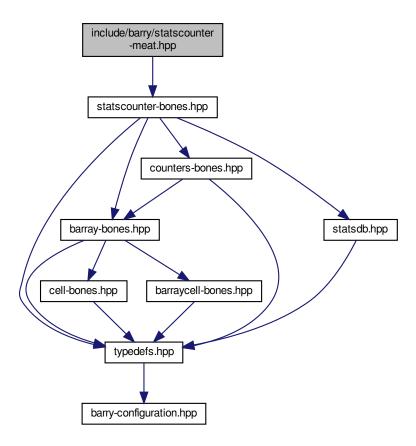
Classes

class StatsCounter< Array_Type, Data_Type >
 Count stats for a single Array.

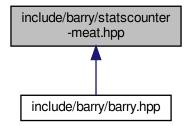
8.42 include/barry/statscounter-meat.hpp File Reference

#include "statscounter-bones.hpp"

Include dependency graph for statscounter-meat.hpp:



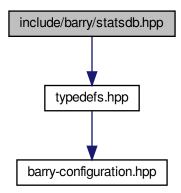
This graph shows which files directly or indirectly include this file:



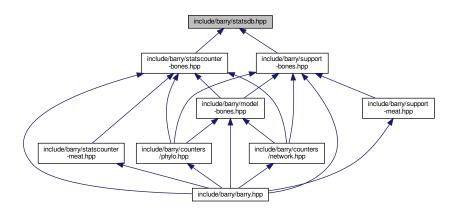
8.43 include/barry/statsdb.hpp File Reference

#include "typedefs.hpp"

Include dependency graph for statsdb.hpp:



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



Classes

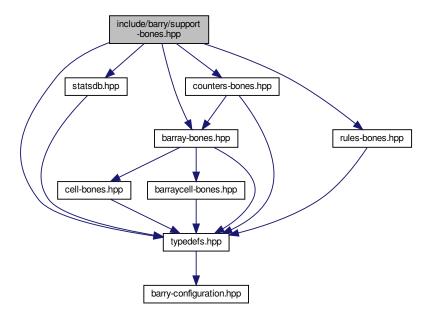
class FreqTable < T >

Database of statistics.

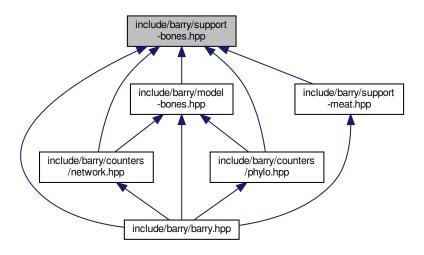
8.44 include/barry/support-bones.hpp File Reference

```
#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:

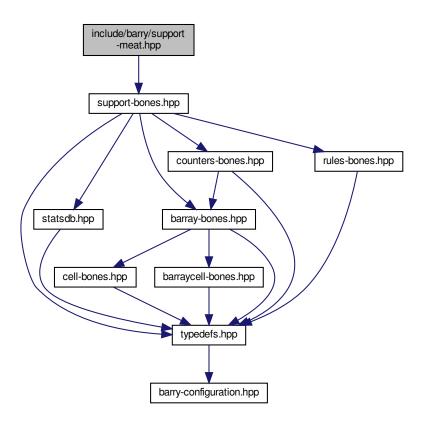


Classes

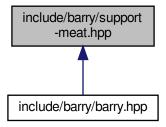
class Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >
 Compute the support of sufficient statistics.

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
- 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

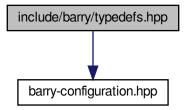
```
std::vector< Array_Type > std::vector< std::vector< double > >* stats_bank

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

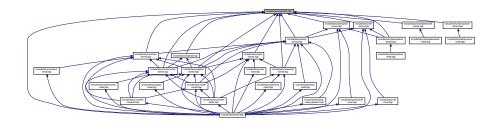
Definition at line 110 of file support-meat.hpp.

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 >, 150
    BArray< Cell Type, Data Type >, 31
\simBArrayCell
                                                     \simStatsCounter
                                                          StatsCounter< Array_Type, Data_Type >, 154
    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 >, 99
    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
                                                              117
    ConstBArrayRowlter< Cell Type, Data Type >, 80
                                                          StatsCounter < Array_Type, Data_Type >, 154
\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 >, 143
\simGeese
                                                         Rules < Array_Type, Data_Type >, 151
    Geese, 104
                                                         Support<
                                                                      Array Type,
                                                                                     Data Counter Type,
\simModel
                                                              Data_Rule_Type, Data_Rule_Dyn_Type >,
    Model<
                Array Type,
                                Data Counter Type,
                                                              160
         add rule dyn
         116
                                                         Model <
                                                                     Array_Type,
                                                                                     Data Counter Type,
\simNetCounterData
                                                              Data_Rule_Type, Data_Rule_Dyn_Type >,
    NetCounterData, 126
                                                              118, 119
\simNetworkData
                                                                      Array_Type,
                                                          Support<
                                                                                     Data_Counter_Type,
    NetworkData, 128
                                                              Data_Rule_Type, Data_Rule_Dyn_Type >,
\simNode
                                                              161
    Node, 131
                                                     annotations
\simPhyloRuleDynData
                                                         Node, 132
    PhyloRuleDynData, 139
                                                     Array
\simPowerSet
                                                          ConstBArrayRowIter< Cell Type, Data Type >, 80
    PowerSet < Array_Type, Data_Rule_Type >, 143
                                                     arrav
\simRule
                                                         Node, 132
    Rule < Array_Type, Data_Type >, 148
                                                     array_bank
```

support-meat.hpp, 232	BARRY_BARRAY_MEAT_OPERATORS_HPP,
arrays	172
Node, 132	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	barray-bones.hpp, 170
PhyloCounterData, 137	BArrayCell
,	BArrayCell< Cell_Type, Data_Type >, 41
BArray	BArrayCell< Cell_Type, Data_Type >, 41
BArray< Cell_Type, Data_Type >, 30, 31	~BArrayCell, 42
BArray< Cell_Type, Data_Type >, 27	BArray< Cell_Type, Data_Type >, 40
∼BArray, 31	BArrayCell, 41
BArray, 30, 31	BArrayDense< Cell_Type, Data_Type >, 61
BArrayCell< Cell Type, Data Type >, 40	
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
— ·	operator=, 43
flush_data, 32	operator==, 43
get_cell, 32	BArrayCell_const
get_col_vec, 32, 33	BArrayCell_const< Cell_Type, Data_Type >, 44
get_entries, 33	BArrayCell_const< Cell_Type, Data_Type >, 43
get_row_vec, 33	\sim BArrayCell_const, 44
insert_cell, 33, 34	BArray < Cell_Type, Data_Type >, 40
is_empty, 34	BArrayCell_const, 44
ncol, 34	BArrayDense < Cell_Type, Data_Type >, 61
nnozero, 34	operator Cell_Type, 44
nrow, 35	operator!=, 45
operator*=, 35	operator<, 45
operator(), 35	operator<=, 45
operator+=, 35, 36	operator>, 45
operator-=, 36	operator>=, 46
operator/=, 36	4-
operator=, 36, 37	operator==, 45
operator==, 37	BArrayDense
out_of_range, 37	BArrayDense < Cell_Type, Data_Type >, 49, 50
print, 37	BArrayDense < Cell_Type, Data_Type >, 46
reserve, 37	~BArrayDense, 50
resize, 37	BArrayCell< Cell_Type, Data_Type >, 61
rm cell, 38	BArrayCell_const< Cell_Type, Data_Type >, 61
- ·	BArrayDense, 49, 50
row, 38	clear, 50
set_data, 38	col, 5 1
swap_cells, 38	D, 51
swap_cols, 39	default_val, 51
swap_rows, 39	get_cell, 51
toggle_cell, 39	get_col_vec, 52
toggle_lock, 39	get_entries, 52
transpose, 39	get_row_vec, 52, 53
visited, 40	insert_cell, 53
zero_col, 40	is_empty, 54
zero_row, 40	ncol, 54
barray-bones.hpp	nnozero, 54
BARRAY_BONES_HPP, 170	nrow, 54
barray-meat-operators.hpp	IIIOw, OT

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: 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 >, 151
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, 137	colnames
PowerSet < Array_Type, Data_Rule_Type >, 143	Flock, 94
blengths	Geese, 105
NodeData, 135	Model
BOTH	Data_Rule_Type, Data_Rule_Dyn_Type >,
CHECK, 24	119
EXISTS, 25	ConstBArrayRowlter
calc	ConstBArrayRowlter < Cell_Type, Data_Type >, 79
PowerSet< Array_Type, Data_Rule_Type >, 144	ConstBArrayRowlter< Cell_Type, Data_Type >, 79
Support< Array_Type, Data_Counter_Type,	~ConstBArrayRowIter, 80
Data_Rule_Type, Data_Rule_Dyn_Type >,	Array, 80
161	ConstBArrayRowlter, 79
calc_backend	current_col, 80
support-meat.hpp, 227	current_row, 80
calc_reduced_sequence	iter, 80
Geese, 104	coordinates_free
calc_sequence	PowerSet < Array_Type, Data_Rule_Type >, 145
Geese, 105	Support< Array_Type, Data_Counter_Type,
Cell	Data_Rule_Type, Data_Rule_Dyn_Type >,
Cell< Cell_Type >, 75, 76	165
Cell< Cell_Type >, 74	coordinates_locked
~Cell, 75	PowerSet < Array_Type, Data_Rule_Type >, 145
add, 76, 77	Support Array_Type, Data_Counter_Type,
Cell, 75, 76	Data_Rule_Type, Data_Rule_Dyn_Type >,
operator Cell_Type, 77	165
operator!=, 77	Count
operator=, 77, 78	Counter< Array_Type, Data_Type >, 83
operator==, 78	count_all
value, 78	StatsCounter< Array_Type, Data_Type >, 155
visited, 78	count_current StateCounter < Array Time Data Time > 155
cfree	StatsCounter< Array_Type, Data_Type >, 155
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 >, 155
164	Counter Counter Array Type Data Type > 92 93
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, <mark>85</mark>	Network counters, 13
name, <mark>85</mark>	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 PhyloPuloPupData 140
counter_istar2	PhyloRuleDynData, 140
Network counters, 12	Counts_type
COUNTER_LAMBDA	typedefs.hpp, 237
barry.hpp, 186	current_col ConstPArroyPoyultor < Coll Type Date Type > 90
counter_longest	ConstBArrayRowlter< Cell_Type, Data_Type >, 80
Phylo counters, 17	current_row ConstBArrayRowIter< Cell_Type, Data_Type >, 80
counter_loss 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	165
	103
Network counters, 13 counter_neofun	D
Phylo counters, 18	BArray< Cell_Type, Data_Type >, 32
counter_neofun_a2b	BArrayDense< Cell_Type, Data_Type >, 51
SSG.MOI_HOOKHI_GED	Rule < Array_Type, Data_Type >, 149

dat	Support< Array_Type, Data_Counter_Type,
Flock, 97	Data_Rule_Type, Data_Rule_Dyn_Type >,
data	161
Counter< Array_Type, Data_Type >, 84	EXISTS, 25
PowerSet< Array_Type, Data_Rule_Type >, 146	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
165	,
support-meat.hpp, 233	f_
delete_data	support-meat.hpp, 233
Counter< Array_Type, Data_Type >, 85	Flock, 92
delete_rengine	∼Flock, 94
Geese, 111	add_data, 94
delete_rules	colnames, 94
Support< Array_Type, Data_Counter_Type,	dat, 97
Data_Rule_Type, Data_Rule_Dyn_Type >,	Flock, 93
165	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
166	likelihood_joint, 95
support-meat.hpp, 233	model, 98
delete_support	nfunctions, 98
Geese, 111	nfuns, 96
desc	nleafs, 96
Counter< Array_Type, Data_Type >, 85	nnodes, 96
directed	nterms, 96
NetworkData, 128	ntrees, 96
duplication	operator(), 96
Node, 133	parse_polytomies, 97
NodeData, 136	rengine, 98
PhyloRuleDynData, 140	set_seed, 97
alaa	support_size, 97
else	flush_data
support-meat.hpp, 233	BArray< Cell_Type, Data_Type >, 32
empty PhyloCounterPate 197	for
PhyloCounterData, 137	support-meat.hpp, 228
EmptyArray PowerSet < Array Type Data Bula Type > 146	FreqTable
PowerSet< Array_Type, Data_Rule_Type >, 146	FreqTable < T >, 99
end PArrow/vector < Cell Type Data Type > 69	FreqTable < T >, 98
BArray Vector < Cell_Type, Data_Type >, 68	~FreqTable, 99
BArrayVector_const< Cell_Type, Data_Type >, 72	add, 99
PhyloCounterData, 137	as_vector, 100
PowerSet < Array_Type, Data_Rule_Type >, 144	clear, 100
Entries Call Time > 01	FreqTable, 99
Entries < Cell_Type >, 91	get_data, 100
Entries Cell_Type >, 90	get_data_ptr, 100
∼Entries, 91	print, 100
Entries, 91	reserve, 101
resize, 91	size, 101
source, 92	Goosa 101
target, 92	Geese, 101
val, 92	~Geese, 104
eval_rules_dyn	calc_reduced_sequence, 104

	calc_sequence, 105 colnames, 105	get_counters Flock, 94
	delete_rengine, 111	Geese, 105
	delete_support, 111	Model < Array_Type, Data_Counter_Type,
	Geese, 103, 104	Data_Rule_Type, Data_Rule_Dyn_Type >,
		119
	get_annotated_nodes, 105	PhyloCounterData, 138
	get_counters, 105	•
	get_model, 105	StatsCounter < Array_Type, Data_Type >, 155
	get_probabilities, 105	Support Array_Type, Data_Counter_Type,
	get_rengine, 106	Data_Rule_Type, Data_Rule_Dyn_Type >,
	get_states, 106	162
	get_support, 106	get_counts
	inherit_support, 106	Support< Array_Type, Data_Counter_Type,
	init, 106	Data_Rule_Type, Data_Rule_Dyn_Type >,
	init_node, 107	162
	initialized, 111	get_counts_ptr
	likelihood, 107	Support< Array_Type, Data_Counter_Type,
	likelihood_exhaust, 107	Data_Rule_Type, Data_Rule_Dyn_Type >,
	map_to_nodes, 111	162
	nfunctions, 111	get_current_stats
	nfuns, 107	Support< Array_Type, Data_Counter_Type,
	nleafs, 107	Data_Rule_Type, Data_Rule_Dyn_Type >,
	nnodes, 108	162
	nodes, 112	get_data
	nterms, 108	FreqTable $<$ T $>$, 100
	observed_counts, 108	PowerSet < Array_Type, Data_Rule_Type >, 144
	operator=, 108	Support< Array_Type, Data_Counter_Type,
	parse_polytomies, 108	$Data_Rule_Type, Data_Rule_Dyn_Type >,$
	predict, 109	162
	predict_backend, 109	get_data_ptr
	predict_exhaust, 109	FreqTable $<$ T $>$, 100
	predict_exhaust_backend, 109	PowerSet< Array_Type, Data_Rule_Type >, 144
	predict_sim, 109	get_entries
	print_observed_counts, 110	BArray< Cell_Type, Data_Type >, 33
	reduced_sequence, 112	BArrayDense < Cell_Type, Data_Type >, 52
	sequence, 112	get_last_name
	set_seed, 110	phylo.hpp, 202
	simulate, 110	get_model
	support_size, 110	Flock, 95
	update_annotations, 110	Geese, 105
gee	se-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	119
	vec_diff, 209	get_parent
	vector_caster, 209	Node, 131
aen	_key	get_probabilities
J	Model< Array_Type, Data_Counter_Type,	Geese, 105
	Data_Rule_Type, Data_Rule_Dyn_Type >,	get_pset
	119	Model< Array_Type, Data_Counter_Type,
aet	annotated_nodes	Data_Rule_Type, Data_Rule_Dyn_Type >,
ອິ່	Geese, 105	120
get_		get_pset_stats
gu_	BArray< Cell_Type, Data_Type >, 32	Model Array_Type, Data_Counter_Type,
	BArrayDense< Cell_Type, Data_Type >, 52	Data_Rule_Type, Data_Rule_Dyn_Type >,
net	col_vec	120
get_	BArray< Cell_Type, Data_Type >, 32, 33	get_rengine
	BArrayDense< Cell_Type, Data_Type >, 52	Geese, 106
	_ Jr - / Jr - / /	,

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
120	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
120	include/barry/models/geese/geese-meat-predict.hpp,
Support< Array_Type, Data_Counter_Type,	212
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/models/geese/geese-meat-predict_exhaust.hpp,
163	212
get_rules_dyn	<pre>include/barry/models/geese/geese-meat-predict_sim.hpp,</pre>
Model< Array_Type, Data_Counter_Type,	213
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/models/geese/geese-meat-simulate.hpp,
120	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,
163	214
get_seq	include/barry/powerset-bones.hpp, 215
Rules < Array_Type, Data_Type >, 151	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, 106	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
121	include/barry/typedefs.hpp, 235
id	indices
Node, 133	NetCounterData, 126
if	inherit_support
support-meat.hpp, 228	Geese, 106
include/barry/barray-bones.hpp, 169	init
	Counter< Array_Type, Data_Type >, 83
include/barry/barray-iterator.hpp, 170	Flock, 95
include/barry/barray-meat-operators.hpp, 171	Geese, 106
include/barry/barray-meat.hpp, 173	init_fun
include/barry/barraycell-bones.hpp, 174	Counter< Array_Type, Data_Type >, 85
include/barry/barraycell-meat.hpp, 175	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 >, 144
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	163
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, 111
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	
include/barry/counters/phylo.hpp, 197	BArrayDense < Cell_Type, Data_Type >, 53
include/barry/model-bones.hpp, 203	support-meat.hpp, 228
include/barry/model-meat.hpp, 204	is_col
include/barry/models/geese.hpp, 206	BArrayVector< Cell_Type, Data_Type >, 68
morado/barry/moders/geese.mpp, 200	

BArrayVector_const< Cell_Type, Data_Type >, 72	add_array, 117
is_empty	add_counter, 117
BArray< Cell_Type, Data_Type >, 34	add_rule, 118
BArrayDense< Cell_Type, Data_Type >, 54	add_rule_dyn, 118, 119
is leaf	colnames, 119
Node, 132	gen_key, 119
is_row	get_counters, 119
BArrayVector< Cell_Type, Data_Type >, 69	-
	get_norm_const, 119
BArrayVector_const< Cell_Type, Data_Type >, 72	get_pset, 120
iter	get_pset_stats, 120
ConstBArrayRowlter< Cell_Type, Data_Type >, 80	get_rengine, 120
koven default	get_rules, 120
keygen_default	get_rules_dyn, 120
model-bones.hpp, 204	get_support, 121
keygen_full	likelihood, 121
geese-bones.hpp, 209	likelihood_total, 121
II-	Model, 116
lb Did Did Did Did 140	nterms, 122
PhyloRuleDynData, 140	operator=, 122
likelihood	print_stats, 122
Geese, 107	sample, 122
Model< Array_Type, Data_Counter_Type,	set_counters, 123
Data_Rule_Type, Data_Rule_Dyn_Type >,	set_keygen, 123
121	set_rengine, 123
likelihood_	set_rules, 123
model-meat.hpp, 205	set_rules_dyn, 123
likelihood_exhaust	set_seed, 124
Geese, 107	size, 124
likelihood_joint	
Flock, 95	size_unique, 124
likelihood_total	store_psets, 124
Model< Array_Type, Data_Counter_Type,	support_size, 124
Data Rule Type, Data Rule Dyn Type >,	model-bones.hpp
121	keygen_default, 204
121	model-meat.hpp
M	likelihood_, 205
PowerSet< Array_Type, Data_Rule_Type >, 146	MODEL_TEMPLATE, 205, 206
Support< Array_Type, Data_Counter_Type,	MODEL_TEMPLATE_ARGS, 205
Data_Rule_Type, Data_Rule_Dyn_Type >,	MODEL_TYPE, 205
166	update_normalizing_constant, 206
Map	MODEL_TEMPLATE
barry-configuration.hpp, 184	model-meat.hpp, 205, 206
	MODEL_TEMPLATE_ARGS
map_to_nodes	model-meat.hpp, 205
Geese, 111	MODEL TYPE
MapVec_type	model-meat.hpp, 205
typedefs.hpp, 237	117
max_num_elements	N
Support< Array_Type, Data_Counter_Type,	PowerSet < Array_Type, Data_Rule_Type >, 146
Data_Rule_Type, Data_Rule_Dyn_Type >,	Support< Array_Type, Data_Counter_Type,
166	Data_Rule_Type, Data_Rule_Dyn_Type >,
Model	166
Model< Array_Type, Data_Counter_Type,	name
Data_Rule_Type, Data_Rule_Dyn_Type >,	Counter< Array_Type, Data_Type >, 85
116	narray
model	Node, 133
Flock, 98	ncol
Model < Array_Type, Data_Counter_Type, Data_Rule_Typ	
Data_Rule_Dyn_Type >, 112	= :: ay < 00::;po; = a:a;po > ; 0 :
~Model, 116	BArrayDense < Cell_Type, Data_Type >, 54
model, 110	NET_C_DATA_IDX

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, 125	network.hpp, 194
∼NetCounterData, 126	NETWORK_COUNTER_LAMBDA
indices, 126	network.hpp, 194
NetCounterData, 125	NETWORK_RULE
numbers, 126	network.hpp, 195
NetCounters	NETWORK_RULE_LAMBDA
network.hpp, 195	network.hpp, 195
NetModel	NetworkData, 126
network.hpp, 196	\sim NetworkData, 128
NetRule	directed, 128
network.hpp, 196	NetworkData, 127, 128
NetRules	vertex_attr, 128
network.hpp, 196	nfunctions
NetStatsCounter	Flock, 98
network.hpp, 196	Geese, 111
NetSupport	nfuns
network.hpp, 196	Flock, 96
Network	Geese, 107
network.hpp, 196	nleafs
Network counters, 10	Flock, 96
counter_absdiff, 11	Geese, 107
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, 129
counter_isolates, 12	~Node, 131
counter_istar2, 12	annotations, 132
counter_mutual, 13	array, 132
counter nodecov, 13	arrays, 132
counter_nodeicov, 13	duplication, 133
counter_nodematch, 13	get_parent, 131
counter nodeocov, 13	id, 133
counter_odegree, 14	is_leaf, 132
counter_odegree15, 14	narray, 133
counter ostar2, 14	Node, 130, 131
counter_ttriads, 14	noffspring, 132
NETWORK_COUNTER, 14	offspring, 133
network.hpp	ord, 133
NET C DATA IDX, 194	parent, 134
NET C DATA NUM, 194	probability, 134
NetCounter, 195	•
	subtree_prob, 134
NetCounters, 195	visited, 134
NetModel, 196	NodeData, 135
NetRule, 196	blengths, 135
NetRules, 196	duplication, 136
NetStatsCounter, 196	NodeData, 135
NetSupport, 196	states, 136
Network, 196	nodes
NETWORK_COUNTER, 194	Geese, 112
NETWORK_COUNTER_LAMBDA, 194	noffspring

Node, 132 NONE	BArrayDenseCell_const< Cell_Type, Data_Type >, 66
CHECK, 24	BArrayVector_const< Cell_Type, Data_Type >, 73
EXISTS, 26	operator*=
nrow	BArray< Cell_Type, Data_Type >, 35
BArray< Cell_Type, Data_Type >, 35	BArrayCell< Cell_Type, Data_Type >, 42
BArrayDense< Cell_Type, Data_Type >, 54	BArrayDense< Cell_Type, Data_Type >, 55
nterms	BArrayDenseCell< Cell_Type, Data_Type >, 63
Flock, 96	BArrayVector< Cell_Type, Data_Type >, 69
Geese, 108	operator()
Model < Array_Type, Data_Counter_Type,	BArray< Cell_Type, Data_Type >, 35
Data_Rule_Type, Data_Rule_Dyn_Type >,	BArrayDense< Cell_Type, Data_Type >, 54, 55
122	Flock, 96
ntrees	PhyloCounterData, 138
Flock, 96	Rule< Array_Type, Data_Type >, 149
numbers	Rules < Array_Type, Data_Type >, 152
NetCounterData, 126	vecHasher< T >, 167
Notobalitor Bata, 120	operator+=
observed counts	BArray< Cell_Type, Data_Type >, 35, 36
Geese, 108	BArrayCell< Cell_Type, Data_Type >, 42
offspring	BArrayDense < Cell_Type, Data_Type >, 55
Node, 133	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 >, 62	BArrayVector< Cell_Type, Data_Type >, 69
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
BArrayVector_const< Cell_Type, Data_Type >, 72	BArrayVector< Cell_Type, Data_Type >, 70 operator=
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37
$\label{eq:barrayVector_const} \mbox{BArrayVector_const} < \mbox{Cell_Type, Data_Type} >, \mbox{72} \\ \mbox{operator!} =$	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayDense< Cell_Type, Data_Type >, 56, 57
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayDense< Cell_Type, Data_Type >, 56, 57 BArrayDenseCell< Cell_Type, Data_Type >, 63
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65 BArrayVector_const< Cell_Type, Data_Type >, 72	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayDense< Cell_Type, Data_Type >, 56, 57 BArrayDenseCell< Cell_Type, Data_Type >, 63 BArrayVector< Cell_Type, Data_Type >, 70
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65 BArrayVector_const< Cell_Type, Data_Type >, 72 Cell< Cell_Type >, 77	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayDense< Cell_Type, Data_Type >, 56, 57 BArrayDenseCell< Cell_Type, Data_Type >, 63 BArrayVector< Cell_Type, Data_Type >, 70 Cell< Cell_Type >, 77, 78
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65 BArrayVector_const< Cell_Type, Data_Type >, 72 Cell< Cell_Type >, 77 operator<	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayDense< Cell_Type, Data_Type >, 56, 57 BArrayDenseCell< Cell_Type, Data_Type >, 63 BArrayVector< Cell_Type, Data_Type >, 70 Cell< Cell_Type >, 77, 78 Counter< Array_Type, Data_Type >, 84
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65 BArrayVector_const< Cell_Type, Data_Type >, 72 Cell< Cell_Type >, 77 operator< BArrayCell_const< Cell_Type, Data_Type >, 45	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayDense< Cell_Type, Data_Type >, 56, 57 BArrayDenseCell< Cell_Type, Data_Type >, 63 BArrayVector< Cell_Type, Data_Type >, 70 Cell< Cell_Type >, 77, 78 Counter< Array_Type, Data_Type >, 84 Counters< Array_Type, Data_Type >, 88, 89
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65 BArrayVector_const< Cell_Type, Data_Type >, 72 Cell< Cell_Type >, 77 operator< BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayDense< Cell_Type, Data_Type >, 56, 57 BArrayDenseCell< Cell_Type, Data_Type >, 63 BArrayVector< Cell_Type, Data_Type >, 70 Cell< Cell_Type >, 77, 78 Counter< Array_Type, Data_Type >, 84 Counters< Array_Type, Data_Type >, 88, 89 Geese, 108
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65 BArrayVector_const< Cell_Type, Data_Type >, 72 Cell< Cell_Type >, 77 operator< BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayDense< Cell_Type, Data_Type >, 56, 57 BArrayDenseCell< Cell_Type, Data_Type >, 63 BArrayVector< Cell_Type, Data_Type >, 70 Cell< Cell_Type >, 77, 78 Counter< Array_Type, Data_Type >, 84 Counters< Array_Type, Data_Type >, 88, 89 Geese, 108 Model< Array_Type, Data_Counter_Type,
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65 BArrayVector_const< Cell_Type, Data_Type >, 72 Cell< Cell_Type >, 77 operator< BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65 BArrayVector_const< Cell_Type, Data_Type >, 73	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayDense< Cell_Type, Data_Type >, 56, 57 BArrayDenseCell< Cell_Type, Data_Type >, 63 BArrayVector< Cell_Type, Data_Type >, 70 Cell< Cell_Type >, 77, 78 Counter< Array_Type, Data_Type >, 84 Counters< Array_Type, Data_Type >, 88, 89 Geese, 108
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65 BArrayVector_const< Cell_Type, Data_Type >, 72 Cell< Cell_Type >, 77 operator< BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65 BArrayVector_const< Cell_Type, Data_Type >, 73 operator<=	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayDense< Cell_Type, Data_Type >, 56, 57 BArrayDenseCell< Cell_Type, Data_Type >, 63 BArrayVector< Cell_Type, Data_Type >, 70 Cell< Cell_Type >, 77, 78 Counter< Array_Type, Data_Type >, 84 Counters< Array_Type, Data_Type >, 88, 89 Geese, 108 Model< Array_Type, Data_Rule_Dyn_Type >, Data_Rule_Type, Data_Rule_Dyn_Type >, 122
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65 BArrayVector_const< Cell_Type, Data_Type >, 72 Cell< Cell_Type >, 77 operator< BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45 BArrayVector_const< Cell_Type, Data_Type >, 65 BArrayVector_const< Cell_Type, Data_Type >, 73 operator<= BArrayCell_const< Cell_Type, Data_Type >, 45	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayDense< Cell_Type, Data_Type >, 56, 57 BArrayDenseCell< Cell_Type, Data_Type >, 63 BArrayVector< Cell_Type, Data_Type >, 70 Cell< Cell_Type >, 77, 78 Counter< Array_Type, Data_Type >, 84 Counters< Array_Type, Data_Type >, 88, 89 Geese, 108 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 122 Rules< Array_Type, Data_Type >, 152
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65 BArrayVector_const< Cell_Type, Data_Type >, 72 Cell< Cell_Type >, 77 operator< BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45 BArrayVector_const< Cell_Type, Data_Type >, 73 operator<= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayCell_const< Cell_Type, Data_Type >, 73 operator<= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayDense< Cell_Type, Data_Type >, 56, 57 BArrayDenseCell< Cell_Type, Data_Type >, 63 BArrayVector< Cell_Type, Data_Type >, 70 Cell< Cell_Type >, 77, 78 Counter< Array_Type, Data_Type >, 84 Counters< Array_Type, Data_Type >, 88, 89 Geese, 108 Model< Array_Type, Data_Rule_Dyn_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 122 Rules< Array_Type, Data_Type >, 152 operator==
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65 BArrayVector_const< Cell_Type, Data_Type >, 72 Cell< Cell_Type >, 77 operator< BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45 BArrayVector_const< Cell_Type, Data_Type >, 73 operator<= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayDense< Cell_Type, Data_Type >, 56, 57 BArrayDenseCell< Cell_Type, Data_Type >, 63 BArrayVector< Cell_Type, Data_Type >, 70 Cell< Cell_Type >, 77, 78 Counter< Array_Type, Data_Type >, 84 Counters< Array_Type, Data_Type >, 88, 89 Geese, 108 Model< Array_Type, Data_Rule_Dyn_Type >, 122 Rules< Array_Type, Data_Type >, 152 operator== BArray< Cell_Type, Data_Type >, 37
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65 BArrayVector_const< Cell_Type, Data_Type >, 72 Cell< Cell_Type >, 77 operator< BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45 BArrayVector_const< Cell_Type, Data_Type >, 73 operator<= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45 BArrayVector_const< Cell_Type, Data_Type >, 73	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayDense< Cell_Type, Data_Type >, 56, 57 BArrayDenseCell< Cell_Type, Data_Type >, 63 BArrayVector< Cell_Type, Data_Type >, 70 Cell< Cell_Type >, 77, 78 Counter< Array_Type, Data_Type >, 84 Counters< Array_Type, Data_Type >, 88, 89 Geese, 108 Model< Array_Type, Data_Rule_Dyn_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 122 Rules< Array_Type, Data_Type >, 152 operator==
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65 BArrayVector_const< Cell_Type, Data_Type >, 72 Cell< Cell_Type >, 77 operator< BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45 BArrayVector_const< Cell_Type, Data_Type >, 73 operator<= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 73 operator<= BArrayDenseCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45 BArrayVector_const< Cell_Type, Data_Type >, 45 BArrayVector_const< Cell_Type, Data_Type >, 73 operator>	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayDense< Cell_Type, Data_Type >, 56, 57 BArrayDenseCell< Cell_Type, Data_Type >, 63 BArrayVector< Cell_Type, Data_Type >, 70 Cell< Cell_Type >, 77, 78 Counter< Array_Type, Data_Type >, 84 Counters< Array_Type, Data_Type >, 88, 89 Geese, 108 Model< Array_Type, Data_Type >, 88, 89 Geese, 108 Model< Array_Type, Data_Rule_Dyn_Type >, 122 Rules< Array_Type, Data_Type >, 152 operator== BArray< Cell_Type, Data_Type >, 37 BArrayCell< Cell_Type, Data_Type >, 43
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65 BArrayVector_const< Cell_Type, Data_Type >, 72 Cell< Cell_Type >, 77 operator< BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45 BArrayVector_const< Cell_Type, Data_Type >, 73 operator<= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 66 BArrayVector_const< Cell_Type, Data_Type >, 73 operator> BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayDense< Cell_Type, Data_Type >, 56, 57 BArrayDenseCell< Cell_Type, Data_Type >, 63 BArrayVector< Cell_Type, Data_Type >, 70 Cell< Cell_Type >, 77, 78 Counter< Array_Type, Data_Type >, 84 Counters< Array_Type, Data_Type >, 88, 89 Geese, 108 Model< Array_Type, Data_Rule_Dyn_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 122 Rules< Array_Type, Data_Type >, 152 operator== BArray< Cell_Type, Data_Type >, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayCell_const< Cell_Type, Data_Type >, 45
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayDense< Cell_Type, Data_Type >, 56, 57 BArrayDenseCell< Cell_Type, Data_Type >, 63 BArrayVector< Cell_Type, Data_Type >, 70 Cell< Cell_Type >, 77, 78 Counter< Array_Type, Data_Type >, 84 Counters< Array_Type, Data_Type >, 88, 89 Geese, 108 Model< Array_Type, Data_Rule_Dyn_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 122 Rules< Array_Type, Data_Type >, 152 operator== BArray< Cell_Type, Data_Type >, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDense< Cell_Type, Data_Type >, 57
BArrayVector_const< Cell_Type, Data_Type >, 72 operator!= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 65 BArrayVector_const< Cell_Type, Data_Type >, 72 Cell< Cell_Type >, 77 operator< BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45 BArrayVector_const< Cell_Type, Data_Type >, 73 operator<= BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 66 BArrayVector_const< Cell_Type, Data_Type >, 73 operator> BArrayCell_const< Cell_Type, Data_Type >, 45 BArrayDenseCell_const< Cell_Type, Data_Type >, 45	BArrayVector< Cell_Type, Data_Type >, 70 operator= BArray< Cell_Type, Data_Type >, 36, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayDense< Cell_Type, Data_Type >, 56, 57 BArrayDenseCell< Cell_Type, Data_Type >, 63 BArrayVector< Cell_Type, Data_Type >, 70 Cell< Cell_Type >, 77, 78 Counter< Array_Type, Data_Type >, 84 Counters< Array_Type, Data_Type >, 88, 89 Geese, 108 Model< Array_Type, Data_Rule_Dyn_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 122 Rules< Array_Type, Data_Type >, 152 operator== BArray< Cell_Type, Data_Type >, 37 BArrayCell< Cell_Type, Data_Type >, 43 BArrayCell< Cell_Type, Data_Type >, 45 BArrayDense< Cell_Type, Data_Type >, 57 BArrayDenseCell< Cell_Type, Data_Type >, 64

BArrayVector_const< Cell_Type, Data_Type >, 73	PhyloCounter
Cell< Cell_Type >, 78	phylo.hpp, 200
operator[]	PhyloCounterData, 136
Counters < Array_Type, Data_Type >, 89	at, 137
PowerSet < Array_Type, Data_Rule_Type >, 145	begin, 137
ord	empty, 137
Node, 133	end, 137
out_of_range	get_counters, 138
BArray< Cell_Type, Data_Type >, 37	operator(), 138
BArrayDense< Cell_Type, Data_Type >, 57	PhyloCounterData, 137
parent	push_back, 138
Node, 134	reserve, 138
parse_polytomies	shrink_to_fit, 138
Flock, 97	size, 138
Geese, 108	PhyloCounters
Phylo counters, 15	phylo.hpp, 200
counter_co_opt, 16	PhyloModel
counter_cogain, 16	phylo.hpp, 201
counter_gains, 16	PhyloPowerSet
	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, 139
counter_overall_changes, 18	\sim PhyloRuleDynData, 139
counter_overall_gains, 19	counts, 140
counter_overall_loss, 19	duplication, 140
counter_prop_genes_changing, 19	lb, 140
counter_subfun, 19	PhyloRuleDynData, 139
Phylo rules, 20	pos, 140
rule_dyn_limit_changes, 20	ub, 140
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	phylo.hpp, 202
PhyloCounter, 200	PhyloSupport
PhyloCounters, 200	phylo.hpp, 202
PhyloModel, 201	POS
PhyloPowerSet, 201	barraydense-meet.hpp, 179
PhyloRule, 201	barraydensecell-meat.hpp, 181
PhyloRuleData, 201	pos
PhyloRuleDyn, 201	PhyloRuleDynData, 140
PhyloRules, 201	PowerSet
PhyloRulesDyn, 202	PowerSet < Array_Type, Data_Rule_Type >, 142
PhyloStatsCounter, 202	PowerSet< Array_Type, Data_Rule_Type >, 141
PhyloSupport, 202	∼PowerSet, 143
PHYLO_CHECK_MISSING	add_rule, 143
phylo.hpp, 199	begin, 143
PHYLO_COUNTER_LAMBDA	calc, 144
phylo.hpp, 199	coordinates_free, 145
PHYLO_RULE_DYN_LAMBDA	coordinates_locked, 145
phylo.hpp, 200	data, 146
PhyloArray	EmptyArray, 146
phylo.hpp, 200	٠٠٠٠ (((((((((((((((((

end, 144	resize
get_data, 144	BArray< Cell_Type, Data_Type >, 37
get_data_ptr, 144	BArrayDense < Cell_Type, Data_Type >, 58
init_support, 144	Entries < Cell_Type >, 91
M, 146	return
N, 146	support-meat.hpp, 234
operator[], 145	rm cell
PowerSet, 142	BArray< Cell_Type, Data_Type >, 38
reset, 145	BArrayDense< Cell Type, Data Type >, 58
rules, 146	support-meat.hpp, 228
rules_deleted, 147	ROW
size, 145	barray-meat-operators.hpp, 172
predict	barray-meat.hpp, 174
Geese, 109	barraydense-meet.hpp, 179
predict_backend	row
Geese, 109	BArray< Cell_Type, Data_Type >, 38
predict_exhaust	BArrayDense< Cell_Type, Data_Type >, 58
Geese, 109	Row_type
predict_exhaust_backend	typedefs.hpp, 237
Geese, 109	Rule
predict_sim	Rule < Array_Type, Data_Type >, 148
Geese, 109	Rule < Array_Type, Data_Type >, 147
	~Rule, 148
print PArroy Coll Type Data Type > 37	~ nule, 146 D, 149
BArray Cell_Type, Data_Type >, 37	•
BArrayDense < Cell_Type, Data_Type >, 57	operator(), 149
FreqTable < T >, 100	Rule, 148
Support< Array_Type, Data_Counter_Type,	rule_dyn_limit_changes
Data_Rule_Type, Data_Rule_Dyn_Type >,	Phylo rules, 20
163	rule_fun_default
print_observed_counts	rules-bones.hpp, 219
Geese, 110	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
122	geese-bones.hpp, 209
printf_barry	RULE_LAMBDA
barry-configuration.hpp, 184	barry.hpp, 187
probability	Rules
Node, 134	Rules < Array_Type, Data_Type >, 150
push_back	rules
PhyloCounterData, 138	PowerSet < Array_Type, Data_Rule_Type >, 146
DEADME and 000	support-meat.hpp, 234
README.md, 239	Rules< Array_Type, Data_Type >, 149
reduced_sequence	\sim Rules, 150
Geese, 112	add_rule, 151
rengine	clear, 151
Flock, 98	get_seq, 151
reserve	operator(), 152
BArray< Cell_Type, Data_Type >, 37	operator=, 152
BArrayDense< Cell_Type, Data_Type >, 57	Rules, 150
FreqTable $<$ T $>$, 101	size, 152
PhyloCounterData, 138	rules-bones.hpp
reset	rule_fun_default, 219
PowerSet < Array_Type, Data_Rule_Type >, 145	rules_
reset_array	support-meat.hpp, 234
StatsCounter< Array_Type, Data_Type >, 155	rules_deleted
Support< Array_Type, Data_Counter_Type,	PowerSet < Array_Type, Data_Rule_Type >, 147
Data_Rule_Type, Data_Rule_Dyn_Type >, 163, 164	rules_dyn

support-meat.hpp, 234 rules_zerodiag network.hpp, 197	Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 124
	PhyloCounterData, 138
sample Model < Array_Type, Data_Counter_Type,	PowerSet < Array_Type, Data_Rule_Type >, 145
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,	Rules < Array_Type, Data_Type >, 152
122	size_unique
sequence	Model < Array_Type, Data_Counter_Type,
Geese, 112	Data_Rule_Type, Data_Rule_Dyn_Type >, 124
set_counters	source
Model< Array_Type, Data_Counter_Type,	Entries < Cell_Type >, 92
Data_Rule_Type, Data_Rule_Dyn_Type >,	states
123	NodeData, 136
StatsCounter< Array_Type, Data_Type >, 156	Statistical Models, 9
Support< Array_Type, Data_Counter_Type,	stats_bank
Data_Rule_Type, Data_Rule_Dyn_Type >,	support-meat.hpp, 234
164	StatsCounter
set_data	StatsCounter< Array_Type, Data_Type >, 154
BArray Cell_Type, Data_Type >, 38	StatsCounter< Array_Type, Data_Type >, 153
BArrayDense < Cell_Type, Data_Type >, 58	~StatsCounter, 154
set_keygen Model< Array Type, Data Counter Type,	add_counter, 154
Data_Rule_Type, Data_Rule_Dyn_Type >,	count_all, 155
123	count_current, 155 count_init, 155
set_rengine	get_counters, 155
Model< Array_Type, Data_Counter_Type,	reset_array, 155
Data_Rule_Type, Data_Rule_Dyn_Type >,	set_counters, 156
123	StatsCounter, 154
set_rules	store_psets
Model< Array_Type, Data_Counter_Type,	Model < Array_Type, Data_Counter_Type,
Data_Rule_Type, Data_Rule_Dyn_Type >,	Data_Rule_Type, Data_Rule_Dyn_Type >,
123	124
Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,	subtree_prob
164	Node, 134
set_rules_dyn	Support Support Array_Type, Data_Counter_Type,
Model< Array_Type, Data_Counter_Type,	Data_Rule_Type, Data_Rule_Dyn_Type >,
Data_Rule_Type, Data_Rule_Dyn_Type >,	159
123	Support < Array_Type, Data_Counter_Type, Data_Rule_Type,
Support< Array_Type, Data_Counter_Type,	Data_Rule_Dyn_Type >, 156
Data_Rule_Type, Data_Rule_Dyn_Type >,	\sim Support, 160
164	add_counter, 160
set_seed	add_rule, 160
Flock, 97	add_rule_dyn, 161
Geese, 110	calc, 161
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,	change_stats, 164
124	coordinates_free, 165
shrink_to_fit	coordinates_locked, 165 current_stats, 165
PhyloCounterData, 138	delete_counters, 165
simulate	delete_rules, 165
Geese, 110	delete_rules_dyn, 166
size	eval_rules_dyn, 161
${\sf BArrayVector}{<}~{\sf Cell_Type},~{\sf Data_Type}>, {\sf 70}$	get_counters, 162
BArrayVector_const< Cell_Type, Data_Type >, 74	get_counts, 162
Counters < Array_Type, Data_Type >, 90	get_counts_ptr, 162
FreqTable $<$ T $>$, 101	get_current_stats, 162

act data 160	torget
get_data, 162	target
get_rules, 163	Entries < Cell_Type >, 92
get_rules_dyn, 163	toggle_cell
init_support, 163	BArray< Cell_Type, Data_Type >, 39
M, 166	BArrayDense < Cell_Type, Data_Type >, 59
max_num_elements, 166	toggle_lock
N, 166	BArray< Cell_Type, Data_Type >, 39
print, 163	BArrayDense< Cell_Type, Data_Type >, 60
reset_array, 163, 164	transpose
set_counters, 164	BArray< Cell_Type, Data_Type >, 39
set_rules, 164	BArrayDense < Cell_Type, Data_Type >, 60
set_rules_dyn, 164	TWO
Support, 159	CHECK, 24
support-meat.hpp	EXISTS, 26
array_bank, 232	typedefs.hpp
BARRY_SUPPORT_MEAT_HPP, 226	Col_type, 237
	Counter_fun_type, 237
calc_backend, 227	
cfree, 232	Counts_type, 237
counters, 232	MapVec_type, 237
counters_, 232	Row_type, 237
delete_counters, 233	Rule_fun_type, 238
delete_rules, 233	uint, 238
delete_rules_dyn, 233	vec_equal, 238
else, 233	vec_equal_approx, 238
f_, 233	vec_inner_prod, 239
for, 228	
if, 228	ub
insert_cell, 228	PhyloRuleDynData, 140
return, 234	uint
rm_cell, 228	typedefs.hpp, 238
rules, 234	UKNOWN
	EXISTS, 26
rules_, 234	update_annotations
rules_dyn, 234	Geese, 110
stats_bank, 234	
SUPPORT_TEMPLATE, 227, 229–232	update_normalizing_constant
SUPPORT_TEMPLATE_ARGS, 227	model-meat.hpp, 206
SUPPORT_TYPE, 227	vol
support_size	val
Flock, 97	Entries < Cell_Type >, 92
Geese, 110	value
Model< Array_Type, Data_Counter_Type,	Cell< Cell_Type >, 78
Data_Rule_Type, Data_Rule_Dyn_Type >,	vec_diff
124	geese-bones.hpp, 209
SUPPORT TEMPLATE	vec_equal
support-meat.hpp, 227, 229–232	typedefs.hpp, 238
SUPPORT_TEMPLATE_ARGS	vec_equal_approx
	typedefs.hpp, 238
support-meat.hpp, 227	vec_inner_prod
SUPPORT_TYPE	typedefs.hpp, 239
support-meat.hpp, 227	vecHasher< T >, 167
swap_cells	
BArray< Cell_Type, Data_Type >, 38	operator(), 167
BArrayDense < Cell_Type, Data_Type >, 59	vector_caster
swap_cols	geese-bones.hpp, 209
BArray< Cell_Type, Data_Type >, 39	vertex_attr
BArrayDense < Cell_Type, Data_Type >, 59	NetworkData, 128
swap rows	visited
BArray< Cell_Type, Data_Type >, 39	BArray< Cell_Type, Data_Type >, 40
BArrayDense < Cell_Type, Data_Type >, 59	BArrayDense < Cell_Type, Data_Type >, 61
a, _ aa 、 a 1, po, _ aa_ 1, po > , a.	Cell< Cell_Type >, 78

```
Node, 134

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
```