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()	17
5.4.2.4 counter_gains_k_offspring()	17
5.4.2.5 counter_genes_changing()	17
5.4.2.6 counter_longest()	17
5.4.2.7 counter_loss()	18
5.4.2.8 counter_maxfuns()	18
5.4.2.9 counter_neofun()	18
5.4.2.10 counter_neofun_a2b()	18
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_subfun()	19
6 Namespace Documentation	2 1
6.1 barry Namespace Reference	21
6.1.1 Detailed Description	
6.2 barry::counters Namespace Reference	
6.2.1 Detailed Description	21
6.3 barry::counters::network Namespace Reference	22
6.4 barry::counters::phylo Namespace Reference	22
6.5 CHECK Namespace Reference	22
6.5.1 Detailed Description	22
6.5.2 Variable Documentation	22
6.5.2.1 BOTH	22
6.5.2.2 NONE	22
6.5.2.3 ONE	22
6.5.2.4 TWO	23
6.6 EXISTS Namespace Reference	23
6.6.1 Detailed Description	23
6.6.2 Variable Documentation	23
6.6.2.1 AS_ONE	23
6.6.2.2 AS_ZERO	23
6.6.2.3 BOTH	
6.6.2.4 NONE	
6.6.2.5 ONE	
6.6.2.6 TWO	
6.6.2.7 UKNOWN	24
7 Class Documentation	25
7.1 BArray< Cell_Type, Data_Type > Class Template Reference	25
7.1.1 Detailed Description	27
7.1.2 Constructor & Destructor Documentation	28
7.1.2.1 BArrav() [1/6]	28

7.1.2.2 BArray() [2/6]	 28
7.1.2.3 BArray() [3/6]	 28
7.1.2.4 BArray() [4/6]	 29
7.1.2.5 BArray() [5/6]	 29
7.1.2.6 BArray() [6/6]	 29
7.1.2.7 ~BArray()	 29
7.1.3 Member Function Documentation	 29
7.1.3.1 clear()	 29
7.1.3.2 col()	 30
7.1.3.3 D() [1/2]	 30
7.1.3.4 D() [2/2]	 30
7.1.3.5 default_val()	 30
7.1.3.6 get_cell()	 30
7.1.3.7 get_col()	 30
7.1.3.8 get_col_vec() [1/2]	 31
7.1.3.9 get_col_vec() [2/2]	 31
7.1.3.10 get_entries()	 31
7.1.3.11 get_row()	 31
7.1.3.12 get_row_vec() [1/2]	 31
7.1.3.13 get_row_vec() [2/2]	 32
7.1.3.14 insert_cell() [1/3]	 32
7.1.3.15 insert_cell() [2/3]	 32
7.1.3.16 insert_cell() [3/3]	 32
7.1.3.17 is_empty()	 32
7.1.3.18 ncol()	 33
7.1.3.19 nnozero()	 33
7.1.3.20 nrow()	 33
7.1.3.21 operator()() [1/2]	 33
7.1.3.22 operator()() [2/2]	 33
7.1.3.23 operator*=()	 33
7.1.3.24 operator+=() [1/3]	 34
7.1.3.25 operator+=() [2/3]	 34
7.1.3.26 operator+=() [3/3]	 34
7.1.3.27 operator-=() [1/3]	 34
7.1.3.28 operator-=() [2/3]	 34
7.1.3.29 operator-=() [3/3]	 34
7.1.3.30 operator/=()	 35
7.1.3.31 operator=() [1/2]	 35
7.1.3.32 operator=() [2/2]	 35
7.1.3.33 operator==()	 35
7.1.3.34 out_of_range()	 35
7.1.3.35 print()	 35

7.1.3.36 reserve()	36
7.1.3.37 resize()	36
7.1.3.38 rm_cell()	36
7.1.3.39 row()	36
7.1.3.40 set_data()	36
7.1.3.41 swap_cells()	37
7.1.3.42 swap_cols()	37
7.1.3.43 swap_rows()	37
7.1.3.44 toggle_cell()	37
7.1.3.45 toggle_lock()	38
7.1.3.46 transpose()	38
7.1.3.47 zero_col()	38
7.1.3.48 zero_row()	38
7.1.4 Friends And Related Function Documentation	38
7.1.4.1 BArrayCell < Cell_Type, Data_Type >	38
7.1.4.2 BArrayCell_const < Cell_Type, Data_Type >	39
7.1.5 Member Data Documentation	39
7.1.5.1 visited	39
7.2 BArrayCell< Cell_Type, Data_Type > Class Template Reference	39
7.2.1 Detailed Description	39
7.2.2 Constructor & Destructor Documentation	40
7.2.2.1 BArrayCell()	40
7.2.2.2 ~BArrayCell()	40
7.2.3 Member Function Documentation	40
7.2.3.1 operator Cell_Type()	40
7.2.3.2 operator*=()	40
7.2.3.3 operator+=()	41
7.2.3.4 operator-=()	41
7.2.3.5 operator/=()	41
7.2.3.6 operator=()	41
7.2.3.7 operator==()	41
7.3 BArrayCell_const< Cell_Type, Data_Type > Class Template Reference	42
7.3.1 Detailed Description	42
7.3.2 Constructor & Destructor Documentation	42
7.3.2.1 BArrayCell_const()	42
7.3.2.2 ~BArrayCell_const()	42
7.3.3 Member Function Documentation	43
7.3.3.1 operator Cell_Type()	43
7.3.3.2 operator"!=()	43
7.3.3.3 operator<()	43
7.3.3.4 operator<=()	43
7.3.3.5 operator==()	43

7.3.3.6 operator>()	 . 44
7.3.3.7 operator>=()	 . 44
7.4 BArrayCol < Cell_Type, Data_Type > Class Template Reference	 . 44
7.4.1 Detailed Description	 . 44
7.4.2 Constructor & Destructor Documentation	 . 45
7.4.2.1 BArrayCol()	 . 45
7.4.2.2 ~BArrayCol()	 . 45
7.4.3 Member Function Documentation	 . 45
7.4.3.1 begin()	 . 45
7.4.3.2 end()	 . 45
7.4.3.3 operator Cell_Type()	 . 46
7.4.3.4 operator*=()	 . 46
7.4.3.5 operator+=()	 . 46
7.4.3.6 operator-=()	 . 46
7.4.3.7 operator/=()	 . 46
7.4.3.8 operator=()	 . 47
7.4.3.9 operator==()	 . 47
7.5 BArrayCol_const< Cell_Type, Data_Type > Class Template Reference	 . 47
7.5.1 Detailed Description	 . 47
7.5.2 Constructor & Destructor Documentation	 . 48
7.5.2.1 ∼BArrayCol_const()	 . 48
7.5.3 Member Function Documentation	 . 48
7.5.3.1 BArrayCol()	 . 48
7.5.3.2 operator"!=()	 . 48
7.5.3.3 operator<()	 . 48
7.5.3.4 operator<=()	 . 49
7.5.3.5 operator==()	 . 49
7.5.3.6 operator>()	 . 49
7.5.3.7 operator>=()	 . 49
7.6 Cell< Cell_Type > Class Template Reference	 . 49
7.6.1 Detailed Description	 . 50
7.6.2 Constructor & Destructor Documentation	 . 50
7.6.2.1 Cell() [1/7]	 . 50
7.6.2.2 Cell() [2/7]	 . 51
7.6.2.3 ~Cell()	 . 51
7.6.2.4 Cell() [3/7]	 . 51
7.6.2.5 Cell() [4/7]	 . 51
7.6.2.6 Cell() [5/7]	 . 51
7.6.2.7 Cell() [6/7]	 . 52
7.6.2.8 Cell() [7/7]	 . 52
7.6.3 Member Function Documentation	 . 52
7.6.3.1 add() [1/4]	 . 52

7.6.3.2 add() [2/4]	52
7.6.3.3 add() [3/4]	52
7.6.3.4 add() [4/4]	53
7.6.3.5 operator Cell_Type()	53
7.6.3.6 operator=() [1/2]	53
7.6.3.7 operator=() [2/2]	53
7.6.4 Member Data Documentation	53
7.6.4.1 value	53
7.6.4.2 visited	54
7.7 ConstBArrayRowlter< Cell_Type, Data_Type > Class Template Reference	54
7.7.1 Detailed Description	55
7.7.2 Constructor & Destructor Documentation	55
7.7.2.1 ConstBArrayRowlter()	55
7.7.2.2 ~ConstBArrayRowlter()	55
7.7.3 Member Data Documentation	55
7.7.3.1 Array	55
7.7.3.2 current_col	55
7.7.3.3 current_row	56
7.7.3.4 iter	56
7.8 Counter< Array_Type, Data_Type > Class Template Reference	56
7.8.1 Detailed Description	57
7.8.2 Constructor & Destructor Documentation	57
7.8.2.1 Counter() [1/3]	57
7.8.2.2 Counter() [2/3]	57
7.8.2.3 Counter() [3/3]	58
7.8.2.4 ~Counter()	58
7.8.3 Member Function Documentation	58
7.8.3.1 count()	58
7.8.3.2 init()	58
7.8.3.3 operator=()	58
7.8.4 Member Data Documentation	59
7.8.4.1 count_fun	59
7.8.4.2 data	59
7.8.4.3 delete_data	59
7.8.4.4 desc	59
7.8.4.5 init_fun	59
7.8.4.6 name	60
$ 7.9 \ Counters < Array_Type, \ Data_Type > Class \ Template \ Reference \\ \ \ldots \\ \ \ldots$	60
7.9.1 Detailed Description	60
7.9.2 Constructor & Destructor Documentation	61
7.9.2.1 Counters() [1/2]	61
7.9.2.2 ~Counters()	61

7.9.2.3 Counters() [2/2]	61
7.9.3 Member Function Documentation	61
7.9.3.1 add_counter() [1/3]	61
7.9.3.2 add_counter() [2/3]	62
7.9.3.3 add_counter() [3/3]	62
7.9.3.4 clear()	62
7.9.3.5 operator=()	62
7.9.3.6 operator[]()	62
7.9.3.7 size()	63
7.10 Entries < Cell_Type > Class Template Reference	63
7.10.1 Detailed Description	63
7.10.2 Constructor & Destructor Documentation	64
7.10.2.1 Entries() [1/2]	64
7.10.2.2 Entries() [2/2]	64
7.10.2.3 ~Entries()	64
7.10.3 Member Function Documentation	64
7.10.3.1 resize()	64
7.10.4 Member Data Documentation	65
7.10.4.1 source	65
7.10.4.2 target	65
7.10.4.3 val	65
7.11 Flock Class Reference	65
7.11.1 Detailed Description	66
7.11.2 Constructor & Destructor Documentation	66
7.11.2.1 Flock()	66
7.11.2.2 ~Flock()	67
7.11.3 Member Function Documentation	67
7.11.3.1 add_data()	67
7.11.3.2 counters_ptr()	67
7.11.3.3 init()	67
7.11.3.4 likelihood_joint()	68
7.11.3.5 nfuns()	68
7.11.3.6 nleafs()	68
7.11.3.7 nnodes()	68
7.11.3.8 nterms()	69
7.11.3.9 ntrees()	69
7.11.3.10 operator()()	69
7.11.3.11 set_seed()	69
7.11.4 Member Data Documentation	70
7.11.4.1 dat	70
7.11.4.2 initialized	70
7.11.4.3 nfunctions	70

7.11.4.4 rengine	70
7.11.4.5 support	70
7.12 FreqTable < T > Class Template Reference	71
7.12.1 Detailed Description	71
7.12.2 Constructor & Destructor Documentation	71
7.12.2.1 FreqTable()	71
7.12.2.2 ~FreqTable()	71
7.12.3 Member Function Documentation	72
7.12.3.1 add()	72
7.12.3.2 as_vector()	72
7.12.3.3 clear()	72
7.12.3.4 get_data()	72
7.12.3.5 get_data_ptr()	72
7.12.3.6 print()	73
7.12.3.7 reserve()	73
7.12.3.8 size()	73
7.13 Geese Class Reference	73
7.13.1 Detailed Description	75
7.13.2 Constructor & Destructor Documentation	75
7.13.2.1 Geese() [1/4]	75
7.13.2.2 Geese() [2/4]	75
7.13.2.3 Geese() [3/4]	76
7.13.2.4 Geese() [4/4]	76
7.13.2.5 ∼Geese()	76
7.13.3 Member Function Documentation	76
7.13.3.1 calc_reduced_sequence()	76
7.13.3.2 calc_sequence()	76
7.13.3.3 get_probabilities()	77
7.13.3.4 inherit_support()	77
7.13.3.5 init()	77
7.13.3.6 init_node()	77
7.13.3.7 likelihood()	77
7.13.3.8 likelihood_exhaust()	78
7.13.3.9 nfuns()	78
7.13.3.10 nleafs()	78
7.13.3.11 nnodes()	78
7.13.3.12 nterms()	78
7.13.3.13 observed_counts()	78
7.13.3.14 operator=() [1/2]	79
7.13.3.15 operator=() [2/2]	79
7.13.3.16 predict()	79
7.13.3.17 predict_backend()	79

/	.13.3.18 print_observed_counts()	79
7	.13.3.19 set_seed()	80
7	.13.3.20 simulate()	80
7	.13.3.21 update_annotations()	80
7.13.4 Me	ember Data Documentation	80
7	.13.4.1 counters	80
7	.13.4.2 delete_counters	80
7	.13.4.3 delete_rengine	81
7	.13.4.4 delete_support	81
7	.13.4.5 initialized	81
7	.13.4.6 map_to_nodes	81
7	.13.4.7 nfunctions	81
7	.13.4.8 nodes	81
7	.13.4.9 reduced_sequence	82
7	.13.4.10 rengine	82
7	.13.4.11 sequence	82
7	.13.4.12 states	82
7	.13.4.13 support	82
	Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class Temerence	83
		86
		86
		86
		86
		87
		87
7.14.3 Me	ember Function Documentation	87
7	.14.3.1 add_array()	87
7	.14.3.2 add_counter() [1/3]	88
7	.14.3.3 add_counter() [2/3]	88
7	.14.3.4 add_counter() [3/3]	88
7	.14.3.5 add_rule() [1/3]	88
7	.14.3.6 add_rule() [2/3]	89
7	.14.3.7 add_rule() [3/3]	89
7	.14.3.8 add_rule_dyn() [1/3]	89
7	.14.3.9 add_rule_dyn() [2/3]	89
7	.14.3.10 add_rule_dyn() [3/3]	90
7	.14.3.11 get_norm_const()	90
7	.14.3.12 get_pset()	90
		90
7	.14.3.14 likelihood() [1/3]	91
7	.14.3.15 likelihood() [2/3]	91

7.14.3.16 likelihood() [3/3]		91
7.14.3.17 likelihood_total()		91
7.14.3.18 nterms()		92
7.14.3.19 operator=()		92
7.14.3.20 print_stats()		92
7.14.3.21 sample() [1/2]		92
7.14.3.22 sample() [2/2]		93
7.14.3.23 set_counters()		93
7.14.3.24 set_keygen()		93
7.14.3.25 set_rengine()		93
7.14.3.26 set_rules()		94
7.14.3.27 set_rules_dyn()		94
7.14.3.28 set_seed()		94
7.14.3.29 size()		94
7.14.3.30 size_unique()		94
7.14.3.31 store_psets()		95
7.14.4 Member Data Documentation		95
7.14.4.1 array_frequency		95
7.14.4.2 arrays2support		95
7.14.4.3 counter_fun		95
7.14.4.4 counters		96
7.14.4.5 delete_rengine		96
7.14.4.6 first_calc_done		96
7.14.4.7 keygen		96
7.14.4.8 keys2support		97
7.14.4.9 n_arrays_per_stats		97
7.14.4.10 normalizing_constants		97
7.14.4.11 params_last		97
7.14.4.12 pset_arrays		98
7.14.4.13 pset_probs		98
7.14.4.14 pset_stats		98
7.14.4.15 rengine		98
7.14.4.16 rules		98
7.14.4.17 rules_dyn		99
7.14.4.18 stats		99
7.14.4.19 support_fun		99
7.14.4.20 target_stats		99
7.14.4.21 with_pset		99
7.15 NetCounterData Class Reference		100
7.15.1 Detailed Description		100
7.15.2 Constructor & Destructor Documentation		100
7.15.2.1 NetCounterData() [1/2]		100

7.15.2.2 NetCounterData() [2/2]
7.15.2.3 ~NetCounterData()
7.15.3 Member Data Documentation
7.15.3.1 indices
7.15.3.2 numbers
7.16 NetworkData Class Reference
7.16.1 Detailed Description
7.16.2 Constructor & Destructor Documentation
7.16.2.1 NetworkData() [1/3]
7.16.2.2 NetworkData() [2/3]
7.16.2.3 NetworkData() [3/3]
7.16.2.4 ~NetworkData()
7.16.3 Member Data Documentation
7.16.3.1 directed
7.16.3.2 vertex_attr
7.17 Node Class Reference
7.17.1 Detailed Description
7.17.2 Constructor & Destructor Documentation
7.17.2.1 Node() [1/5]
7.17.2.2 Node() [2/5]
7.17.2.3 Node() [3/5]
7.17.2.4 Node() [4/5]
7.17.2.5 Node() [5/5]
7.17.2.6 ~Node()
7.17.3 Member Function Documentation
7.17.3.1 get_parent()
7.17.3.2 is_leaf()
7.17.4 Member Data Documentation
7.17.4.1 annotations
7.17.4.2 array
7.17.4.3 arrays
7.17.4.4 duplication
7.17.4.5 id
7.17.4.6 narray
7.17.4.7 offspring
7.17.4.8 ord
7.17.4.9 parent
7.17.4.10 probability
7.17.4.11 subtree_prob
7.17.4.12 visited
7.18 NodeData Class Reference
7.18.1 Detailed Description 10

7.18.2 Constructor & Destructor Documentation)9
7.18.2.1 NodeData() [1/2])9
7.18.2.2 NodeData() [2/2])9
7.18.2.3 ~NodeData()	0
7.18.3 Member Data Documentation	0
7.18.3.1 blengths	0
7.18.3.2 duplication	0
7.18.3.3 states	0
7.19 PowerSet< Array_Type, Data_Rule_Type > Class Template Reference	1
7.19.1 Detailed Description	2
7.19.2 Constructor & Destructor Documentation	2
7.19.2.1 PowerSet() [1/3]	2
7.19.2.2 PowerSet() [2/3]	2
7.19.2.3 PowerSet() [3/3]	3
7.19.2.4 ~PowerSet()	3
7.19.3 Member Function Documentation	3
7.19.3.1 add_rule() [1/3]11	3
7.19.3.2 add_rule() [2/3]11	3
7.19.3.3 add_rule() [3/3]11	3
7.19.3.4 begin()	4
7.19.3.5 calc()	4
7.19.3.6 end()	4
7.19.3.7 get_data()	4
7.19.3.8 get_data_ptr()	4
7.19.3.9 init_support()	5
7.19.3.10 operator[]()	5
7.19.3.11 reset()	5
7.19.3.12 size()	5
7.19.4 Member Data Documentation	5
7.19.4.1 coordinates_free	5
7.19.4.2 coordinates_locked	6
7.19.4.3 data	6
7.19.4.4 EmptyArray	6
7.19.4.5 M	6
7.19.4.6 N	6
7.19.4.7 rules	7
7.19.4.8 rules_deleted	7
7.20 Rule < Array_Type, Data_Type > Class Template Reference	7
7.20.1 Detailed Description	8
7.20.2 Constructor & Destructor Documentation	8
7.20.2.1 Rule() [1/2]	8
7.20.2.2 Rule() [2/2]	8

7.20.2.3 ~Rule()	8
7.20.3 Member Function Documentation	9
7.20.3.1 D()	9
7.20.3.2 operator()()	9
7.21 Rules < Array_Type, Data_Type > Class Template Reference	9
7.21.1 Detailed Description	:0
7.21.2 Constructor & Destructor Documentation	:0
7.21.2.1 Rules() [1/2]	:0
7.21.2.2 Rules() [2/2]	0
7.21.2.3 ~Rules()	1:1
7.21.3 Member Function Documentation	1
7.21.3.1 add_rule() [1/3]12	1
7.21.3.2 add_rule() [2/3]12	1:1
7.21.3.3 add_rule() [3/3]12	1
7.21.3.4 clear()	1
7.21.3.5 get_seq()	1!
7.21.3.6 operator()()	2
7.21.3.7 operator=()	2
7.21.3.8 size()	23
7.22 StatsCounter< Array_Type, Data_Type > Class Template Reference	23
7.22.1 Detailed Description	<u>'</u> 4
7.22.2 Constructor & Destructor Documentation	4
7.22.2.1 StatsCounter() [1/2]	4
7.22.2.2 StatsCounter() [2/2]	4
7.22.2.3 ~StatsCounter()	:5
7.22.3 Member Function Documentation	:5
7.22.3.1 add_counter() [1/2]	:5
7.22.3.2 add_counter() [2/2]	:5
7.22.3.3 count_all()	:5
7.22.3.4 count_current()	:5
7.22.3.5 count_init()	6
7.22.3.6 reset_array()	6
7.22.3.7 set_counters()	6
7.22.4 Member Data Documentation	6
7.22.4.1 Array	6
7.22.4.2 counter_deleted	27
7.22.4.3 counters	27
7.22.4.4 current_stats	:7
7.22.4.5 EmptyArray	:7
7.23 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class Template Reference	28
7.23.1 Detailed Description	30

7.23.2 Constructor & Destructor Documentation	130
7.23.2.1 Support() [1/3]	130
7.23.2.2 Support() [2/3]	130
7.23.2.3 Support() [3/3]	131
7.23.2.4 ~Support()	131
7.23.3 Member Function Documentation	131
7.23.3.1 add_counter() [1/2]	131
7.23.3.2 add_counter() [2/2]	131
7.23.3.3 add_rule() [1/2]	132
7.23.3.4 add_rule() [2/2]	132
7.23.3.5 add_rule_dyn() [1/2]	132
7.23.3.6 add_rule_dyn() [2/2]	132
7.23.3.7 calc()	132
7.23.3.8 get_counts()	133
7.23.3.9 get_counts_ptr()	133
7.23.3.10 get_current_stats()	133
7.23.3.11 init_support()	133
7.23.3.12 print()	134
7.23.3.13 reset_array() [1/2]	134
7.23.3.14 reset_array() [2/2]	134
7.23.3.15 set_counters()	134
7.23.3.16 set_rules()	134
7.23.3.17 set_rules_dyn()	135
7.23.4 Member Data Documentation	135
7.23.4.1 change_stats	135
7.23.4.2 coordinates_free	135
7.23.4.3 coordinates_locked	135
7.23.4.4 counters	136
7.23.4.5 current_stats	136
7.23.4.6 data	136
7.23.4.7 delete_counters	136
7.23.4.8 delete_rules	137
7.23.4.9 delete_rules_dyn	137
7.23.4.10 EmptyArray	137
7.23.4.11 M	137
7.23.4.12 max_num_elements	138
7.23.4.13 N	138
7.23.4.14 rules	138
7.23.4.15 rules_dyn	138
7.24 vecHasher < T > Struct Template Reference	139
7.24.1 Detailed Description	139
7.24.2 Member Function Documentation	139

	7.24.2.1 operator()()	139
8	File Documentation	141
	8.1 include/barry/barray-bones.hpp File Reference	141
	8.1.1 Macro Definition Documentation	142
	8.1.1.1 BARRAY_BONES_HPP	142
	8.2 include/barry/barray-iterator.hpp File Reference	142
	8.2.1 Macro Definition Documentation	142
	8.2.1.1 BARRAY_ITERATOR_HPP	143
	8.3 include/barry/barray-meat-operators.hpp File Reference	143
	8.3.1 Macro Definition Documentation	144
	8.3.1.1 COL	144
	8.3.1.2 ROW	144
	8.3.2 Function Documentation	144
	8.3.2.1 checkdim_()	144
	8.4 include/barry/barray-meat.hpp File Reference	144
	8.4.1 Macro Definition Documentation	145
	8.4.1.1 COL	145
	8.4.1.2 ROW	145
	8.5 include/barry/barraycell-bones.hpp File Reference	146
	8.6 include/barry/barraycell-meat.hpp File Reference	146
	8.7 include/barry/barraycol-bones.hpp File Reference	147
	8.8 include/barry/barraycol-meat.hpp File Reference	148
	8.8.1 Macro Definition Documentation	148
	8.8.1.1 BARRY_BARRAYCOL_MEAT_HPP	148
	8.9 include/barry/barry-configuration.hpp File Reference	148
	8.9.1 Macro Definition Documentation	149
	8.9.1.1 BARRY_CHECK_SUPPORT	149
	8.9.1.2 BARRY_ISFINITE	149
	8.9.1.3 BARRY_MAX_NUM_ELEMENTS	150
	8.9.1.4 BARRY_SAFE_EXP	150
	8.9.2 Typedef Documentation	150
	8.9.2.1 Map	150
	8.10 include/barry/barry.hpp File Reference	150
	8.10.1 Macro Definition Documentation	151
	8.10.1.1 COUNTER_FUNCTION	151
	8.10.1.2 COUNTER_LAMBDA	152
	8.10.1.3 RULE_FUNCTION	152
	8.10.1.4 RULE_LAMBDA	152
	8.11 include/barry/cell-bones.hpp File Reference	152
	8.12 include/barry/cell-meat.hpp File Reference	153
	8.13 include/barry/col-bones.hpp File Reference	154

8.18.1 Function Documentation	38
8.18.1.1 keygen_default()	38
8.18.1.2 likelihood_()	38
8.18.1.3 update_normalizing_constant()	36
8.19 include/barry/model-meat.hpp File Reference	39
8.20 include/barry/models/geese.hpp File Reference	39
8.21 include/barry/models/geese/flock-bones.hpp File Reference	70
8.22 include/barry/models/geese/flock-meet.hpp File Reference	70
8.23 include/barry/models/geese/geese-bones.hpp File Reference	71
8.23.1 Macro Definition Documentation	71
8.23.1.1 INITIALIZED	71
8.23.2 Function Documentation	72
8.23.2.1 keygen_full()	72
8.23.2.2 RULE_FUNCTION()	72
8.23.2.3 vec_diff()	72
8.23.2.4 vector_caster()	72
8.24 include/barry/models/geese/geese-meat-constructors.hpp File Reference	73
8.25 include/barry/models/geese/geese-meat-likelihood.hpp File Reference	73
8.26 include/barry/models/geese/geese-meat-likelihood_exhaust.hpp File Reference	74
8.27 include/barry/models/geese/geese-meat-predict.hpp File Reference	75
8.28 include/barry/models/geese/geese-meat-simulate.hpp File Reference	75
8.29 include/barry/models/geese/geese-meat.hpp File Reference	76
8.30 include/barry/models/geese/geese-node-bones.hpp File Reference	77
8.31 include/barry/powerset-bones.hpp File Reference	77
8.32 include/barry/powerset-meat.hpp File Reference	78
8.33 include/barry/rules-bones.hpp File Reference	79
8.33.1 Function Documentation	30
8.33.1.1 rule_fun_default()	30
8.34 include/barry/rules-meat.hpp File Reference	30
8.35 include/barry/statscounter-bones.hpp File Reference	31
8.36 include/barry/statscounter-meat.hpp File Reference	32
8.37 include/barry/statsdb.hpp File Reference	33
8.38 include/barry/support-bones.hpp File Reference	34
8.39 include/barry/support-meat.hpp File Reference	35
8.39.1 Macro Definition Documentation	35
8.39.1.1 BARRY_SUPPORT_MEAT_HPP	35
8.40 include/barry/typedefs.hpp File Reference	36
8.40.1 Typedef Documentation	37
8.40.1.1 Col_type	38
8.40.1.2 Counter_fun_type	38
8.40.1.3 Counts_type	38
8.40.1.4 MapVec_type	38

ndex		191
8.41	ADME.md File Reference	. 190
	8.40.2.3 vec_inner_prod()	. 190
	8.40.2.2 vec_equal_approx()	. 190
	8.40.2.1 vec_equal()	. 189
	40.2 Function Documentation	. 189
	8.40.1.7 uint	. 189
	8.40.1.6 Rule_fun_type	. 189
	8.40.1.5 Row_type	. 189

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();
  std::cout «
    "Edges : " « counts[0] « std::endl «
"Transitive triads : " « counts[1] « std::endl «
"Isolates : " « counts[2] « std::endl «
"C triads : " « counts[3] « std::endl «
"Mutuals : " « counts[4] « std::endl;
 return 0;
Compiling this program using g++
g++ -std=c++11 -Wall -pedantic 08-counts.cpp -o counts && ./counts
Yields the following output:
Current view
                1
  0,] 1 1
   1,] .
            1
  . 1
   5,]
New view
  0,] .
1,] 1
            1 1
  1,] 1 .
2,] 1 .
                    . 1
   3,] .
4,] 1
           . 1 . .
   5,] . . . .
Edges
Transitive triads : 3
Isolates
C triads
```

Code of Conduct

Mutuals

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						 											 					ç
Statistical Models						 											 					ç
Network counters						 											 					10
Phylo counters						 											 					15

4 Module Index

Class Index

3.1 Class List

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

BArray< Ceil_Type, Data_Type >
Baseline class for binary arrays
BArrayCell < Cell_Type, Data_Type >
BArrayCell_const< Cell_Type, Data_Type >
BArrayCol < Cell_Type, Data_Type >
BArrayCol_const< Cell_Type, Data_Type >
Cell< Cell_Type >
Entries in BArray. For now, it only has two members:
ConstBArrayRowlter< Cell_Type, Data_Type >
Counter< Array_Type, Data_Type >
A counter function based on change statistics
Counters < Array_Type, Data_Type >
Vector of counters
Entries < Cell_Type >
A wrapper class to store source, target, val from a BArray object 63
Flock
A Flock is a group of Geese
FreqTable < T >
Database of statistics
Geese
Annotated Phylo Model
Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >
General framework for discrete exponential models. This class allows generating discrete expo-
nential models in the form of a linear exponential model:
NetCounterData
Data class used to store arbitrary uint or double vectors
NetworkData Date class for Networks
Data class for Networks
Node A single node for the model
NodeData
Data definition for the PhyloArray class
PowerSet < Array_Type, Data_Rule_Type >
Powerset of a binary array
Rule < Array_Type, Data_Type >
Rule for determining if a cell should be included in a sequence
Table for determining it a contention of moladed in a coquence

6 Class Index

Rules < Array_Type, Data_Type >	
Vector of objects of class Rule	119
StatsCounter< Array_Type, Data_Type >	
Count stats for a single Array	123
Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >	
Compute the support of sufficient statistics	128
vecHasher< T >	139

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/barraycol-bones.hpp
include/barry/barraycol-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
include/barry/models/geese/flock-meet.hpp
include/barry/models/geese/geese-bones.hpp
include/barry/models/geese/geese-meat-constructors.hpp
include/barry/models/geese/geese-meat-likelihood.hpp

8 File Index

include/barry/models/geese/geese-meat-likelihood_exhaust.hpp	174
include/barry/models/geese/geese-meat-predict.hpp	175
include/barry/models/geese/geese-meat-simulate.hpp	175
include/barry/models/geese/geese-meat.hpp	176
include/barry/models/geese/geese-node-bones.hpp	177

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.

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.

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.

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_overall_loss (PhyloCounters *counters, bool duplication=true)

Overall functional loss.

- void counter_maxfuns (PhyloCounters *counters, uint lb, uint ub, bool duplication=true)
 - Cap the number of functions per gene.
- void counter loss (PhyloCounters *counters, std::vector< uint > nfun, bool duplication=true)

Total count of losses for an specific function.

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

Total count of Sub-functionalization events.

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

Co-evolution (joint gain or loss)

- void counter_longest (PhyloCounters *counters)
 - Longest branch mutates (either by gain or by loss)
- void counter_neofun (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)

Total number of neofunctionalization events.

• void counter_neofun_a2b (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)

Total number of neofunctionalization events.

void counter_co_opt (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)
 Function co-opting.

5.4.1 Detailed Description

Counters for phylogenetic modeling.

Parameters

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

5.4.2 Function Documentation

5.4.2.1 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 1081 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 711 of file phylo.hpp.

5.4 Phylo counters

5.4.2.3 counter_gains()

Functional gains for a specific function (nfun).

Definition at line 149 of file phylo.hpp.

5.4.2.4 counter_gains_k_offspring()

k genes gain function nfun

Definition at line 191 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 265 of file phylo.hpp.

5.4.2.6 counter_longest()

Longest branch mutates (either by gain or by loss)

Definition at line 770 of file phylo.hpp.

18 Module Documentation

5.4.2.7 counter_loss()

Total count of losses for an specific function.

Definition at line 512 of file phylo.hpp.

5.4.2.8 counter_maxfuns()

Cap the number of functions per gene.

Definition at line 428 of file phylo.hpp.

5.4.2.9 counter_neofun()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 875 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 960 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 559 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 109 of file phylo.hpp.

5.4.2.13 counter_overall_loss()

Overall functional loss.

Definition at line 382 of file phylo.hpp.

5.4.2.14 counter_subfun()

Total count of Sub-functionalization events.

It requires to specify data = {funA, funB}

Definition at line 625 of file phylo.hpp.

20 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 30 of file typedefs.hpp.

6.5.2.2 NONE

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

Definition at line 31 of file typedefs.hpp.

6.5.2.3 ONE

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

Definition at line 32 of file typedefs.hpp.

6.5.2.4 TWO

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

Definition at line 33 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 48 of file typedefs.hpp.

6.6.2.2 AS_ZERO

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

Definition at line 47 of file typedefs.hpp.

6.6.2.3 BOTH

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

Definition at line 41 of file typedefs.hpp.

6.6.2.4 NONE

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

Definition at line 42 of file typedefs.hpp.

6.6.2.5 ONE

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

Definition at line 43 of file typedefs.hpp.

6.6.2.6 TWO

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

Definition at line 44 of file typedefs.hpp.

6.6.2.7 UKNOWN

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

Definition at line 46 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
- const Row_type< Cell_Type > * get_row (uint i, bool check_bounds=true) const
- const Col type< Cell Type > * get col (uint i, 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_
catarget h	NWhen 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

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()

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

7.1.3.3 D() [1/2]

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

7.1.3.4 D() [2/2]

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

7.1.3.5 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 get_cell()

7.1.3.7 get_col()

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()

7.1.3.12 get_row_vec() [1/2]

7.1.3.13 get_row_vec() [2/2]

7.1.3.14 insert_cell() [1/3]

7.1.3.15 insert_cell() [2/3]

7.1.3.16 insert_cell() [3/3]

7.1.3.17 is_empty()

7.1.3.18 ncol()

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

7.1.3.19 nnozero()

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

7.1.3.20 nrow()

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

7.1.3.21 operator()() [1/2]

7.1.3.22 operator()() [2/2]

7.1.3.23 operator*=()

7.1.3.24 operator+=() [1/3]

7.1.3.25 operator+=() [2/3]

7.1.3.26 operator+=() [3/3]

7.1.3.27 operator-=() [1/3]

7.1.3.28 operator-=() [2/3]

7.1.3.29 operator-=() [3/3]

7.1.3.30 operator/=()

7.1.3.31 operator=() [1/2]

Move assignment.

7.1.3.32 operator=() [2/2]

Assignment constructor.

7.1.3.33 operator==()

7.1.3.34 out_of_range()

7.1.3.35 print()

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

7.1.3.36 reserve()

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

7.1.3.37 resize()

7.1.3.38 rm_cell()

7.1.3.39 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
```

7.1.3.40 set_data()

Set the data object.

Parameters

data_	
delete_ <i>←</i>	
data_	

7.1.3.41 swap_cells()

7.1.3.42 swap_cols()

7.1.3.43 swap_rows()

7.1.3.44 toggle_cell()

7.1.3.45 toggle_lock()

7.1.3.46 transpose()

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

7.1.3.47 zero_col()

7.1.3.48 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
- bool operator> (const Cell Type &val) const
- bool operator<= (const Cell_Type &val) const
- bool operator>= (const Cell_Type &val) const

7.3.1 Detailed Description

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

Definition at line 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 ~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 BArrayCol < Cell_Type, Data_Type > Class Template Reference

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

Public Member Functions

```
    BArrayCol (BArray < Cell_Type, Data_Type > *Array_, uint i_, bool check_bounds=true)
```

- ∼BArrayCol ()
- 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
- Col_type< Cell_Type >::iterator begin () noexcept
- Col_type< Cell_Type >::iterator end () noexcept

7.4.1 Detailed Description

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

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

7.4.2 Constructor & Destructor Documentation

7.4.2.1 BArrayCol()

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

7.4.2.2 ~BArrayCol()

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

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

7.4.3 Member Function Documentation

7.4.3.1 begin()

```
template<typename Cell_Type , typename Data_Type >
Col_type< Cell_Type >::iterator BArrayCol< Cell_Type, Data_Type >::begin [inline], [noexcept]
```

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

7.4.3.2 end()

```
template<typename Cell_Type , typename Data_Type >
Col_type< Cell_Type >::iterator BArrayCol< Cell_Type, Data_Type >::end [inline], [noexcept]
```

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

7.4.3.3 operator Cell_Type()

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

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

7.4.3.4 operator*=()

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

7.4.3.5 operator+=()

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

7.4.3.6 operator-=()

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

7.4.3.7 operator/=()

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

7.4.3.8 operator=()

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

7.4.3.9 operator==()

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

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

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

7.5 BArrayCol_const< Cell_Type, Data_Type > Class Template Reference

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

Public Member Functions

- BArrayCol (const BArray < Cell_Type, Data_Type > *Array_, uint i_, bool check_bounds=true)
- ∼BArrayCol_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

7.5.1 Detailed Description

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

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

7.5.2 Constructor & Destructor Documentation

7.5.2.1 \sim BArrayCol_const()

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

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

7.5.3 Member Function Documentation

7.5.3.1 BArrayCol()

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

7.5.3.2 operator"!=()

Definition at line 92 of file barraycol-meat.hpp.

7.5.3.3 operator<()

Definition at line 97 of file barraycol-meat.hpp.

7.5.3.4 operator<=()

Definition at line 107 of file barraycol-meat.hpp.

7.5.3.5 operator==()

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

7.5.3.6 operator>()

Definition at line 102 of file barraycol-meat.hpp.

7.5.3.7 operator>=()

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

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

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

7.6 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
- void add (double x)
- void add (unsigned int x)
- void add (int x)
- Cell ()
- Cell ()
- Cell ()

Public Attributes

- Cell_Type value
- bool visited

7.6.1 Detailed Description

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

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

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

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

7.6.2 Constructor & Destructor Documentation

7.6.2.1 Cell() [1/7]

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

7.6.2.2 Cell() [2/7]

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

7.6.2.3 ∼CeII()

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

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

7.6.2.4 Cell() [3/7]

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

7.6.2.5 Cell() [4/7]

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

7.6.2.6 Cell() [5/7]

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

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

7.6.2.7 Cell() [6/7]

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

Definition at line 45 of file cell-meat.hpp.

7.6.2.8 Cell() [7/7]

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

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

7.6.3 Member Function Documentation

7.6.3.1 add() [1/4]

7.6.3.2 add() [2/4]

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

7.6.3.3 add() [3/4]

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

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

7.6.3.4 add() [4/4]

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

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

7.6.3.5 operator Cell_Type()

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

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

7.6.3.6 operator=() [1/2]

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

7.6.3.7 operator=() [2/2]

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

7.6.4 Member Data Documentation

7.6.4.1 value

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

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

7.6.4.2 visited

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

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

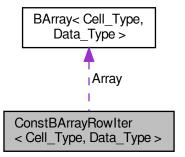
The documentation for this class was generated from the following files:

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

7.7 ConstBArrayRowlter< Cell_Type, Data_Type > Class Template Reference

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

Collaboration diagram for ConstBArrayRowIter< 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

7.7.1 Detailed Description

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

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

7.7.2 Constructor & Destructor Documentation

7.7.2.1 ConstBArrayRowIter()

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

7.7.2.2 ∼ConstBArrayRowlter()

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

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

7.7.3 Member Data Documentation

7.7.3.1 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.

7.7.3.2 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.

7.7.3.3 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.

7.7.3.4 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

7.8 Counter< Array_Type, Data_Type > Class Template Reference

A counter function based on change statistics.

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

Public Member Functions

- Counter< Array_Type, Data_Type > operator= (const Counter< Array_Type, Data_Type > &counter_)
- ∼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_←	When true, the destructor will delete the pointer in the main data.
data_	

- 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_)
```

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 = ""

7.8.1 Detailed Description

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

A counter function based on change statistics.

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

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

7.8.2 Constructor & Destructor Documentation

7.8.2.1 Counter() [1/3]

```
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.

7.8.2.2 Counter() [2/3]

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

7.8.2.3 Counter() [3/3]

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

7.8.2.4 ~Counter()

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

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

7.8.3 Member Function Documentation

7.8.3.1 count()

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

7.8.3.2 init()

Definition at line 136 of file counters-meat.hpp.

7.8.3.3 operator=()

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

7.8.4 Member Data Documentation

7.8.4.1 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.

7.8.4.2 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.

7.8.4.3 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.

7.8.4.4 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.

7.8.4.5 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.

7.8.4.6 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

7.9 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_)
- Counters < Array_Type, Data_Type > operator= (const Counters < Array_Type, Data_Type > &counter_)
- 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 ()

7.9.1 Detailed Description

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

Vector of counters.

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

Definition at line 98 of file counters-bones.hpp.

7.9.2 Constructor & Destructor Documentation

7.9.2.1 Counters() [1/2]

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

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

7.9.2.2 ~Counters()

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

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

7.9.2.3 Counters() [2/2]

Definition at line 67 of file counters-meat.hpp.

7.9.3 Member Function Documentation

7.9.3.1 add_counter() [1/3]

Definition at line 157 of file counters-meat.hpp.

7.9.3.2 add_counter() [2/3]

Definition at line 169 of file counters-meat.hpp.

7.9.3.3 add_counter() [3/3]

Definition at line 180 of file counters-meat.hpp.

7.9.3.4 clear()

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

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

7.9.3.5 operator=()

Definition at line 94 of file counters-meat.hpp.

7.9.3.6 operator[]()

Returns a pointer to a particular counter.

Parameters

```
idx Id of the counter
```

Returns

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

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

7.9.3.7 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 130 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

7.10 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

7.10.1 Detailed Description

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

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

Template Parameters

```
Cell_Type Any type
```

Definition at line 69 of file typedefs.hpp.

7.10.2 Constructor & Destructor Documentation

7.10.2.1 Entries() [1/2]

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

Definition at line 75 of file typedefs.hpp.

7.10.2.2 Entries() [2/2]

Definition at line 76 of file typedefs.hpp.

7.10.2.3 ∼Entries()

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

Definition at line 83 of file typedefs.hpp.

7.10.3 Member Function Documentation

7.10.3.1 resize()

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

Definition at line 85 of file typedefs.hpp.

7.11 Flock Class Reference 65

7.10.4 Member Data Documentation

7.10.4.1 source

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

Definition at line 71 of file typedefs.hpp.

7.10.4.2 target

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

Definition at line 72 of file typedefs.hpp.

7.10.4.3 val

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

Definition at line 73 of file typedefs.hpp.

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

• include/barry/typedefs.hpp

7.11 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 ()
- phylocounters::PhyloCounters * counters_ptr ()

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

Public Attributes

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

7.11.1 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.

7.11.2 Constructor & Destructor Documentation

7.11.2.1 Flock()

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

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

7.11 Flock Class Reference 67

7.11.2.2 ∼Flock()

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

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

7.11.3 Member Function Documentation

7.11.3.1 add_data()

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

Add a tree to the flock.

Parameters

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

Returns

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

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

7.11.3.2 counters_ptr()

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

Definition at line 75 of file flock-meet.hpp.

7.11.3.3 init()

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

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

7.11.3.4 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 84 of file flock-meet.hpp.

7.11.3.5 nfuns()

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

Definition at line 109 of file flock-meet.hpp.

7.11.3.6 nleafs()

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

Definition at line 132 of file flock-meet.hpp.

7.11.3.7 nnodes()

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

Definition at line 121 of file flock-meet.hpp.

7.11 Flock Class Reference 69

7.11.3.8 nterms()

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

Definition at line 144 of file flock-meet.hpp.

7.11.3.9 ntrees()

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

Definition at line 115 of file flock-meet.hpp.

7.11.3.10 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 151 of file flock-meet.hpp.

7.11.3.11 set_seed()

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

Set the seed of the model.

Parameters

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

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

7.11.4 Member Data Documentation

7.11.4.1 dat

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

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

7.11.4.2 initialized

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

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

7.11.4.3 nfunctions

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

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

7.11.4.4 rengine

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

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

7.11.4.5 support

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

Definition at line 23 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

7.12 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

7.12.1 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.

7.12.2 Constructor & Destructor Documentation

7.12.2.1 FreqTable()

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

Definition at line 28 of file statsdb.hpp.

7.12.2.2 ∼FreqTable()

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

Definition at line 29 of file statsdb.hpp.

7.12.3 Member Function Documentation

7.12.3.1 add()

Definition at line 47 of file statsdb.hpp.

7.12.3.2 as_vector()

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

Definition at line 61 of file statsdb.hpp.

7.12.3.3 clear()

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

Definition at line 83 of file statsdb.hpp.

7.12.3.4 get_data()

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

Definition at line 73 of file statsdb.hpp.

7.12.3.5 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.

7.12.3.6 print()

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

Definition at line 102 of file statsdb.hpp.

7.12.3.7 reserve()

Definition at line 89 of file statsdb.hpp.

7.12.3.8 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

7.13 Geese Class Reference

Annotated Phylo Model.

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

Public Member Functions

- ∼Geese ()
- void init ()
- 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)

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 \mathbb{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

- · unsigned int nfuns () const noexcept
- unsigned int nnodes () const noexcept
- unsigned int nleafs () const noexcept
- unsigned int nterms () const

Geese prediction

Calculate the conditional probability

Parameters

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

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

Returns

std::vector< double > Returns the posterior probability

- std::vector< std::vector< double > > predict_backend (const std::vector< double > &par, bool use_
 reduced_sequence, const std::vector< uint > &preorder)

Public Attributes

- unsigned int nfunctions
- std::map< unsigned int, Node > nodes

7.13 Geese Class Reference 75

- 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_counters = false
- bool delete_support = false

Shared objects within a <tt>Geese</tt>

Since users may start adding counters before initializing the PhyloModel object, the object counter is initialized first.

While the member support has an rengine, since Geese can sample trees, we have the option to keep it separate.

```
• std::mt19937 * rengine = nullptr
```

- phylocounters::PhyloCounters * counters = nullptr
- phylocounters::PhyloModel * support = nullptr
- std::vector< std::vector< bool >> states

7.13.1 Detailed Description

Annotated Phylo Model.

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

Definition at line 70 of file geese-bones.hpp.

7.13.2 Constructor & Destructor Documentation

7.13.2.1 Geese() [1/4]

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

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

7.13.2.2 Geese() [2/4]

Definition at line 17 of file geese-meat-constructors.hpp.

7.13.2.3 Geese() [3/4]

Definition at line 157 of file geese-meat-constructors.hpp.

7.13.2.4 Geese() [4/4]

Definition at line 232 of file geese-meat-constructors.hpp.

7.13.2.5 ∼Geese()

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

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

7.13.3 Member Function Documentation

7.13.3.1 calc_reduced_sequence()

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

Definition at line 234 of file geese-meat.hpp.

7.13.3.2 calc_sequence()

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

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

7.13 Geese Class Reference 77

7.13.3.3 get_probabilities()

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

Definition at line 277 of file geese-meat.hpp.

7.13.3.4 inherit_support()

Definition at line 140 of file geese-meat.hpp.

7.13.3.5 init()

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

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

7.13.3.6 init_node()

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

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

7.13.3.7 likelihood()

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

7.13.3.8 likelihood_exhaust()

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

7.13.3.9 nfuns()

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

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

7.13.3.10 nleafs()

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

Definition at line 301 of file geese-meat.hpp.

7.13.3.11 nnodes()

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

Definition at line 297 of file geese-meat.hpp.

7.13.3.12 nterms()

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

Definition at line 311 of file geese-meat.hpp.

7.13.3.13 observed_counts()

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

Definition at line 318 of file geese-meat.hpp.

7.13.3.14 operator=() [1/2]

7.13.3.15 operator=() [2/2]

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

7.13.3.16 predict()

Definition at line 166 of file geese-meat-predict.hpp.

7.13.3.17 predict_backend()

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

7.13.3.18 print_observed_counts()

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

Definition at line 366 of file geese-meat.hpp.

7.13.3.19 set_seed()

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

7.13.3.20 simulate()

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

Definition at line 12 of file geese-meat-simulate.hpp.

7.13.3.21 update_annotations()

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

Definition at line 168 of file geese-meat.hpp.

7.13.4 Member Data Documentation

7.13.4.1 counters

```
phylocounters::PhyloCounters* Geese::counters = nullptr
```

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

7.13.4.2 delete_counters

```
bool Geese::delete_counters = false
```

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

7.13.4.3 delete_rengine

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

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

7.13.4.4 delete_support

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

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

7.13.4.5 initialized

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

Definition at line 100 of file geese-bones.hpp.

7.13.4.6 map_to_nodes

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

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

7.13.4.7 nfunctions

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

Definition at line 91 of file geese-bones.hpp.

7.13.4.8 nodes

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

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

7.13.4.9 reduced_sequence

```
\verb|std::vector<| unsigned int > Geese::reduced_sequence|
```

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

7.13.4.10 rengine

```
std::mt19937* Geese::rengine = nullptr
```

Definition at line 84 of file geese-bones.hpp.

7.13.4.11 sequence

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

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

7.13.4.12 states

```
std::vector< std::vector< bool > > Geese::states
```

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

7.13.4.13 support

```
phylocounters::PhyloModel* Geese::support = nullptr
```

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

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

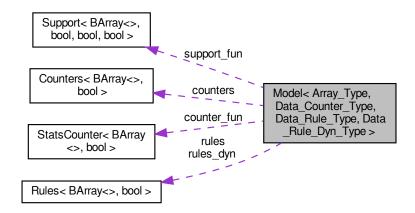
- include/barry/models/geese/geese-bones.hpp
- include/barry/models/geese/geese-meat-constructors.hpp
- include/barry/models/geese/geese-meat-likelihood.hpp
- include/barry/models/geese/geese-meat-likelihood_exhaust.hpp
- include/barry/models/geese/geese-meat-predict.hpp
- include/barry/models/geese/geese-meat-simulate.hpp
- include/barry/models/geese/geese-meat.hpp

7.14 Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class Template Reference

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

#include <model-bones.hpp>

Collaboration diagram for Model < Array Type, Data Counter Type, Data Rule Type, Data Rule Dyn Type >:



Public Member Functions

- 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)
- 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)

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 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)
- $\bullet \ \ \text{void} \ \ \overset{\cdot}{\text{set_rules}} \ (\text{Rules} < \text{Array_Type}, \ \text{Data_Rule_Type} > * \text{rules_)} \\$
- void add_rule_dyn (Rule< Array_Type, Data_Rule_Dyn_Type > &rule)
- void add_rule_dyn (Rule < Array_Type, Data_Rule_Dyn_Type > *rule)
- void set_rules_dyn (Rules < Array_Type, Data_Rule_Dyn_Type > *rules_)

Likelihood functions.

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

Parameters

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

- double likelihood (const std::vector< double > ¶ms, const uint &i, bool as_log=false)
- double likelihood (const std::vector< double > ¶ms, const std::vector< double > &target_, const uint &i, bool as_log=false)
- double likelihood_total (const std::vector< double > ¶ms, bool as_log=false)

Extract elements by index

Parameters

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

- double get norm const (const std::vector< double > ¶ms, const uint &i, bool as log=false)
- const std::vector< Array_Type > * get_pset (const uint &i)
- const std::vector< std::vector< double >> * get_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

Public Attributes

- std::vector< Counts type > stats
- std::vector< uint > n_arrays_per_stats
- MapVec_type< double, uint > keys2support

Map of types of arrays to support sets.

std::vector< std::vector< double >> params_last

Vector of the previously used parameters.

- std::vector< double > normalizing constants
- std::vector< bool > first_calc_done
- std::function< std::vector< double >const Array Type &)> keygen = nullptr

Function to extract features of the array to be hash.

Container space for the powerset (and its sufficient stats)

This is useful in the case of using simulations or evaluating functions that need to account for the full set of states.

- bool with_pset = false
- std::vector< std::vector< Array_Type >> pset_arrays
- std::vector< std::vector< double >>> pset stats
- std::vector< std::vector< double >> pset_probs

Information about the arrays used in the model

target_stats holds the observed sufficient statistics for each array in the dataset. array_frequency contains the frequency with which each of the target stats (arrays) shows in the support. array2support maps array indices (0, 1, ...) to the corresponding support.

- std::vector< std::vector< double >> target_stats
- std::vector< uint > array frequency
- std::vector< uint > arrays2support

Functions to compute statistics

Arguments are recycled to save memory and computation.

- Counters < Array_Type, Data_Counter_Type > counters
- Rules < Array Type, Data Rule Type > rules
- Rules
 Array_Type, Data_Rule_Dyn_Type > rules_dyn
- Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > support_fun
- StatsCounter< Array_Type, Data_Counter_Type > counter_fun

Random number generation

Random number generation

- std::mt19937 * rengine = nullptr
- bool delete rengine = false
- void set_rengine (std::mt19937 *rengine_, bool delete_=false)
- void set_seed (unsigned int s)

7.14.1 Detailed Description

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

class Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >

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

$$\frac{\exp\left(\theta^{\mathsf{t}}c(A)\right)}{\sum_{A'\in\mathcal{A}}\exp\left(\theta^{\mathsf{t}}c(A')\right)}$$

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

Template Parameters

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

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

7.14.2 Constructor & Destructor Documentation

7.14.2.1 Model() [1/3]

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

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

7.14.2.2 Model() [2/3]

Definition at line 27 of file model-meat.hpp.

7.14.2.3 Model() [3/3]

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

7.14.2.4 ∼Model()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\(\circ\)

Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>

Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::\(\circ\)Model () [inline]
```

Definition at line 200 of file model-bones.hpp.

7.14.3 Member Function Documentation

7.14.3.1 add_array()

Adds an array to the support of not already included.

Parameters

Array_	array to be added	
force_new	If false, it will use keygen to obtain a double vector and create a hash of it. If the hash has	
	been computed earlier, the support is recycled.	

Returns

The number of the array.

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

Definition at line 286 of file model-meat.hpp.

7.14.3.2 add_counter() [1/3]

Definition at line 134 of file model-meat.hpp.

7.14.3.3 add_counter() [2/3]

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

7.14.3.4 add counter() [3/3]

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

7.14.3.5 add_rule() [1/3]

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

7.14.3.6 add_rule() [2/3]

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

7.14.3.7 add_rule() [3/3]

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

7.14.3.8 add_rule_dyn() [1/3]

Definition at line 236 of file model-meat.hpp.

7.14.3.9 add_rule_dyn() [2/3]

Definition at line 245 of file model-meat.hpp.

7.14.3.10 add_rule_dyn() [3/3]

Definition at line 255 of file model-meat.hpp.

7.14.3.11 get_norm_const()

Definition at line 517 of file model-meat.hpp.

7.14.3.12 get_pset()

Definition at line 549 of file model-meat.hpp.

7.14.3.13 get_stats()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type > const std::vector< std::vector< double > > * Model< Array_Type, Data_Counter_Type, Data_\leftarrow Rule_Type, Data_Rule_Dyn_Type >::get_stats ( const uint & i ) [inline]
```

Definition at line 562 of file model-meat.hpp.

7.14.3.14 likelihood() [1/3]

Definition at line 403 of file model-meat.hpp.

7.14.3.15 likelihood() [2/3]

Definition at line 443 of file model-meat.hpp.

7.14.3.16 likelihood() [3/3]

Definition at line 370 of file model-meat.hpp.

7.14.3.17 likelihood_total()

Definition at line 477 of file model-meat.hpp.

7.14.3.18 nterms()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
uint Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::nterms [inline],
[noexcept]
```

Definition at line 603 of file model-meat.hpp.

7.14.3.19 operator=()

Definition at line 82 of file model-meat.hpp.

7.14.3.20 print_stats()

Definition at line 574 of file model-meat.hpp.

7.14.3.21 sample() [1/2]

7.14.3.22 sample() [2/2]

Definition at line 610 of file model-meat.hpp.

7.14.3.23 set counters()

Definition at line 172 of file model-meat.hpp.

7.14.3.24 set_keygen()

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

7.14.3.25 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 115 of file model-bones.hpp.

7.14.3.26 set_rules()

Definition at line 223 of file model-meat.hpp.

7.14.3.27 set_rules_dyn()

Definition at line 272 of file model-meat.hpp.

7.14.3.28 set_seed()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\longleftrightarrow Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::set_seed ( unsigned int s ) [inline]
```

Definition at line 125 of file model-bones.hpp.

7.14.3.29 size()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
uint Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::size [inline],
[noexcept]
```

Definition at line 593 of file model-meat.hpp.

7.14.3.30 size_unique()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
uint Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::size_unique
[inline], [noexcept]
```

Definition at line 598 of file model-meat.hpp.

7.14.3.31 store_psets()

template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::store_psets
[inline], [noexcept]

Definition at line 118 of file model-meat.hpp.

7.14.4 Member Data Documentation

7.14.4.1 array frequency

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

Definition at line 163 of file model-bones.hpp.

7.14.4.2 arrays2support

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< uint > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type
>::arrays2support
```

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

7.14.4.3 counter fun

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
StatsCounter<Array_Type, Data_Counter_Type> Model< Array_Type, Data_Counter_Type, Data_Rule_←
Type, Data_Rule_Dyn_Type >::counter_fun
```

Definition at line 182 of file model-bones.hpp.

7.14.4.4 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> Model< Array_Type, Data_Counter_Type, Data_Rule_Dyn_Type >::counters

Definition at line 178 of file model-bones.hpp.

7.14.4.5 delete_rengine

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::delete_←
rengine = false
```

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

7.14.4.6 first calc done

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_← Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> std::vector< bool > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::first_calc_done
```

Definition at line 188 of file model-bones.hpp.

7.14.4.7 keygen

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::function<std::vector<double>const Array_Type &)> Model< Array_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::keygen = nullptr
```

Function to extract features of the array to be hash.

Definition at line 192 of file model-bones.hpp.

7.14.4.8 keys2support

template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>

MapVec_type< double, uint > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule\times
_Dyn_Type >::keys2support

Map of types of arrays to support sets.

This is of the same length as the vector stats.

Definition at line 171 of file model-bones.hpp.

7.14.4.9 n arrays per stats

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

Definition at line 140 of file model-bones.hpp.

7.14.4.10 normalizing_constants

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< double > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_←
Type >::normalizing_constants
```

Definition at line 187 of file model-bones.hpp.

7.14.4.11 params_last

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

Vector of the previously used parameters.

Definition at line 186 of file model-bones.hpp.

7.14.4.12 pset_arrays

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector< Array_Type >> Model< Array_Type, Data_Counter_Type, Data_Rule_←
Type, Data_Rule_Dyn_Type >::pset_arrays
```

Definition at line 149 of file model-bones.hpp.

7.14.4.13 pset probs

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

Definition at line 151 of file model-bones.hpp.

7.14.4.14 pset stats

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

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

7.14.4.15 rengine

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_← Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::mt19937* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >←
::rengine = nullptr
```

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

7.14.4.16 rules

```
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_Type> Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_\times
_Rule_Dyn_Type >::rules
```

Definition at line 179 of file model-bones.hpp.

7.14.4.17 rules dyn

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Rules<Array_Type, Data_Rule_Dyn_Type> Model< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::rules_dyn
```

Definition at line 180 of file model-bones.hpp.

7.14.4.18 stats

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< Counts_type > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_\times
Dyn_Type >::stats
```

Definition at line 139 of file model-bones.hpp.

7.14.4.19 support_fun

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

Definition at line 181 of file model-bones.hpp.

7.14.4.20 target stats

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

Definition at line 162 of file model-bones.hpp.

7.14.4.21 with_pset

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarray_Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
bool Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::with_pset = false
```

Definition at line 148 of file model-bones.hpp.

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

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

7.15 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

7.15.1 Detailed Description

Data class used to store arbitrary uint or double vectors.

Definition at line 61 of file network.hpp.

7.15.2 Constructor & Destructor Documentation

7.15.2.1 NetCounterData() [1/2]

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

Definition at line 67 of file network.hpp.

7.15.2.2 NetCounterData() [2/2]

Definition at line 68 of file network.hpp.

7.15.2.3 ∼NetCounterData()

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

Definition at line 73 of file network.hpp.

7.15.3 Member Data Documentation

7.15.3.1 indices

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

Definition at line 64 of file network.hpp.

7.15.3.2 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

7.16 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

7.16.1 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.

7.16.2 Constructor & Destructor Documentation

7.16.2.1 NetworkData() [1/3]

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

Definition at line 30 of file network.hpp.

7.16.2.2 NetworkData() [2/3]

Constructor using a single attribute.

Parameters

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

Definition at line 38 of file network.hpp.

7.16.2.3 NetworkData() [3/3]

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

Constructor using multiple attributes.

7.17 Node Class Reference 103

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.

7.16.2.4 ~NetworkData()

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

Definition at line 56 of file network.hpp.

7.16.3 Member Data Documentation

7.16.3.1 directed

bool NetworkData::directed = true

Definition at line 27 of file network.hpp.

7.16.3.2 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

7.17 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
- · 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.

7.17.1 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.

7.17.2 Constructor & Destructor Documentation

7.17 Node Class Reference 105

7.17.2.1 Node() [1/5]

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

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

7.17.2.2 Node() [2/5]

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

Definition at line 55 of file geese-node-bones.hpp.

7.17.2.3 Node() [3/5]

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

Definition at line 61 of file geese-node-bones.hpp.

7.17.2.4 Node() [4/5]

```
Node::Node (
          Node && x ) [inline], [noexcept]
```

Definition at line 68 of file geese-node-bones.hpp.

7.17.2.5 Node() [5/5]

```
Node::Node ( {\tt const\ Node\ \&\ x\ )} \quad [{\tt inline}]
```

Definition at line 82 of file geese-node-bones.hpp.

7.17.2.6 ∼Node()

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

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

7.17.3 Member Function Documentation

7.17.3.1 get_parent()

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

Definition at line 96 of file geese-node-bones.hpp.

7.17.3.2 is_leaf()

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

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

7.17.4 Member Data Documentation

7.17.4.1 annotations

```
\verb|std::vector<| unsigned int > \verb|Node::annotations| \\
```

Observed annotations (only defined for Geese)

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

7.17.4.2 array

phylocounters::PhyloArray Node::array

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

7.17 Node Class Reference 107

7.17.4.3 arrays

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

Arrays given all possible states.

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

7.17.4.4 duplication

```
bool Node::duplication
```

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

7.17.4.5 id

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

Id of the node (as specified in the input)

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

7.17.4.6 narray

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

ID of the array in the model.

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

7.17.4.7 offspring

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

Offspring nodes.

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

7.17.4.8 ord

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

Order in which the node was created.

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

7.17.4.9 parent

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

Parent node.

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

7.17.4.10 probability

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

The probability of observing each state.

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

7.17.4.11 subtree_prob

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

Induced subtree probabilities.

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

7.17.4.12 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

7.18 NodeData Class Reference

Data definition for the PhyloArray class.

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

Public Member Functions

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

Public Attributes

- std::vector< double > blengths
- std::vector< bool > states
- bool duplication = true

7.18.1 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.

7.18.2 Constructor & Destructor Documentation

7.18.2.1 NodeData() [1/2]

```
NodeData::NodeData ( ) [inline]
```

Definition at line 41 of file phylo.hpp.

7.18.2.2 NodeData() [2/2]

Definition at line 43 of file phylo.hpp.

7.18.2.3 ∼NodeData()

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

Definition at line 49 of file phylo.hpp.

7.18.3 Member Data Documentation

7.18.3.1 blengths

```
std::vector< double > NodeData::blengths
```

Branch length.

Definition at line 29 of file phylo.hpp.

7.18.3.2 duplication

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

Definition at line 39 of file phylo.hpp.

7.18.3.3 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

7.19 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

7.19.1 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.

7.19.2 Constructor & Destructor Documentation

7.19.2.1 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.

7.19.2.2 PowerSet() [2/3]

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

7.19.2.3 PowerSet() [3/3]

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

7.19.2.4 ∼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.

7.19.3 Member Function Documentation

7.19.3.1 add_rule() [1/3]

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

7.19.3.2 add_rule() [2/3]

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

7.19.3.3 add_rule() [3/3]

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

7.19.3.4 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.

7.19.3.5 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.

7.19.3.6 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.

7.19.3.7 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.

7.19.3.8 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.

7.19.3.9 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.

7.19.3.10 operator[]()

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

7.19.3.11 reset()

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

7.19.3.12 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.

7.19.4 Member Data Documentation

7.19.4.1 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.

7.19.4.2 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.

7.19.4.3 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.

7.19.4.4 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.

7.19.4.5 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.

7.19.4.6 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.

7.19.4.7 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.

7.19.4.8 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

7.20 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)

7.20.1 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.

7.20.2 Constructor & Destructor Documentation

7.20.2.1 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.

7.20.2.2 Rule() [2/2]

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

7.20.2.3 ∼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.

7.20.3 Member Function Documentation

7.20.3.1 D()

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

Read/Write access to the data.

7.20.3.2 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

7.21 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)

7.21.1 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.

7.21.2 Constructor & Destructor Documentation

7.21.2.1 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.

7.21.2.2 Rules() [2/2]

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

7.21.2.3 ∼Rules()

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

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

7.21.3 Member Function Documentation

7.21.3.1 add_rule() [1/3]

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

7.21.3.2 add_rule() [2/3]

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

7.21.3.3 add_rule() [3/3]

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

7.21.3.4 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.

7.21.3.5 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.

7.21.3.6 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.

7.21.3.7 operator=()

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

7.21.3.8 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:

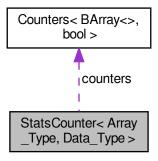
- include/barry/rules-bones.hpp
- include/barry/rules-meat.hpp

7.22 StatsCounter< Array_Type, Data_Type > Class Template Reference

Count stats for a single Array.

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

Collaboration diagram for StatsCounter< Array Type, Data Type >:



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 ()

Public Attributes

- const Array_Type * Array
- Array_Type EmptyArray
- std::vector< double > current_stats
- Counters < Array_Type, Data_Type > * counters
- bool counter_deleted = false

7.22.1 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.

7.22.2 Constructor & Destructor Documentation

7.22.2.1 StatsCounter() [1/2]

Creator of a StatsCounter

Parameters

Array←	A const pointer to a BArray.

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

7.22.2.2 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 49 of file statscounter-bones.hpp.

7.22.2.3 ∼StatsCounter()

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

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

7.22.3 Member Function Documentation

7.22.3.1 add_counter() [1/2]

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

7.22.3.2 add_counter() [2/2]

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

7.22.3.3 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.

7.22.3.4 count_current()

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

7.22.3.5 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.

7.22.3.6 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.

7.22.3.7 set_counters()

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

7.22.4 Member Data Documentation

7.22.4.1 Array

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
const Array_Type* StatsCounter< Array_Type, Data_Type >::Array
```

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

7.22.4.2 counter_deleted

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
bool StatsCounter< Array_Type, Data_Type >::counter_deleted = false
```

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

7.22.4.3 counters

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Counters<Array_Type,Data_Type>* StatsCounter< Array_Type, Data_Type >::counters
```

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

7.22.4.4 current_stats

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
std::vector< double > StatsCounter< Array_Type, Data_Type >::current_stats
```

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

7.22.4.5 EmptyArray

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

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

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

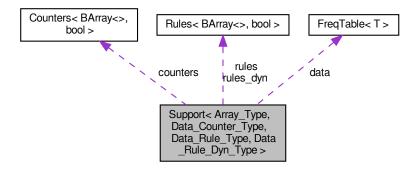
- include/barry/statscounter-bones.hpp
- include/barry/statscounter-meat.hpp

7.23 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>

Collaboration diagram for Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >:



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
- const std::vector< double > & get_current_stats () const

List current statistics.

• void print () const

Resets the support calculator

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

Parameters

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

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

Manage counters

Parameters

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

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

Manage rules

Parameters

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

- void add_rule (Rule < Array_Type, Data_Rule_Type > *f_)
- void add_rule (Rule < Array_Type, Data_Rule_Type > f_)
- void set_rules (Rules < Array_Type, Data_Rule_Type > *rules_)
- void add_rule_dyn (Rule< Array_Type, Data_Rule_Dyn_Type > *f_)
 void add_rule_dyn (Rule< Array_Type, Data_Rule_Dyn_Type > f_)
- void set_rules_dyn (Rules < Array_Type, Data_Rule_Dyn_Type > *rules_)

Public Attributes

Array_Type EmptyArray

Reference array to generate the support.

FreqTable data

Table with the support.

Counters < Array_Type, Data_Counter_Type > * counters

Vector of couter functions.

• Rules< Array Type, Data Rule Type > * rules

Vector of static rules (cells to iterate).

Rules
 Array_Type, Data_Rule_Dyn_Type > * rules_dyn

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

- uint N
- uint M
- bool delete counters = true
- bool delete rules = true
- bool delete_rules_dyn = true
- uint max_num_elements = BARRY_MAX_NUM_ELEMENTS
- std::vector< double > current stats
- std::vector< std::pair< uint, uint >> coordinates_free
- std::vector< std::pair< uint, uint >> coordinates_locked
- std::vector< std::vector< double > > change_stats

7.23.1 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.

7.23.2 Constructor & Destructor Documentation

7.23.2.1 Support() [1/3]

Constructor passing a reference Array.

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

7.23.2.2 Support() [2/3]

Constructor specifying the dimensions of the array (empty).

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

7.23.2.3 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.

7.23.2.4 ~Support()

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

7.23.3 Member Function Documentation

7.23.3.1 add_counter() [1/2]

Definition at line 204 of file support-meat.hpp.

7.23.3.2 add_counter() [2/2]

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

7.23.3.3 add_rule() [1/2]

Definition at line 241 of file support-meat.hpp.

7.23.3.4 add_rule() [2/2]

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

7.23.3.5 add_rule_dyn() [1/2]

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

7.23.3.6 add_rule_dyn() [2/2]

Definition at line 286 of file support-meat.hpp.

7.23.3.7 calc()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::calc (
    std::vector< Array_Type > * array_bank = nullptr,
    std::vector< std::vector< double > > * stats_bank = nullptr,
    unsigned int max_num_elements_ = 0u ) [inline]
```

Computes the entire support.

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

Parameters

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

Definition at line 178 of file support-meat.hpp.

7.23.3.8 get counts()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
Counts_type Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >
::get_counts [inline]
```

Definition at line 313 of file support-meat.hpp.

7.23.3.9 get_counts_ptr()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
const MapVec_type * Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type
>::get_counts_ptr [inline]
```

Definition at line 320 of file support-meat.hpp.

7.23.3.10 get_current_stats()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type > const std::vector< double > & Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_\leftarrow Rule_Dyn_Type >::get_current_stats [inline]
```

List current statistics.

Definition at line 327 of file support-meat.hpp.

7.23.3.11 init_support()

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

7.23.3.12 print()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::print
[inline]
```

Definition at line 332 of file support-meat.hpp.

7.23.3.13 reset_array() [1/2]

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::reset_array
[inline]
```

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

7.23.3.14 reset_array() [2/2]

Definition at line 80 of file support-meat.hpp.

7.23.3.15 set counters()

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

7.23.3.16 set_rules()

Definition at line 261 of file support-meat.hpp.

7.23.3.17 set_rules_dyn()

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

void Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::set_rules←
_dyn (

Rules< Array_Type, Data_Rule_Dyn_Type > * rules_ ) [inline]
```

Definition at line 296 of file support-meat.hpp.

7.23.4 Member Data Documentation

7.23.4.1 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.

7.23.4.2 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.

7.23.4.3 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.

7.23.4.4 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 >::counters

Vector of couter functions.

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

7.23.4.5 current stats

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

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

7.23.4.6 data

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

Table with the support.

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

7.23.4.7 delete_counters

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

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

7.23.4.8 delete_rules

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

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

7.23.4.9 delete_rules_dyn

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

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

7.23.4.10 EmptyArray

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
Array_Type Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >\times
::EmptyArray
```

Reference array to generate the support.

Temp array used to iterate through the support.

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

7.23.4.11 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.

7.23.4.12 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.

7.23.4.13 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.

7.23.4.14 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 >::rules
```

Vector of static rules (cells to iterate).

Definition at line 52 of file support-bones.hpp.

7.23.4.15 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 >::rules_dyn
```

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

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

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

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

7.24 vecHasher < T > Struct Template Reference

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

Public Member Functions

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

7.24.1 Detailed Description

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

Definition at line 96 of file typedefs.hpp.

7.24.2 Member Function Documentation

7.24.2.1 operator()()

Definition at line 97 of file typedefs.hpp.

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

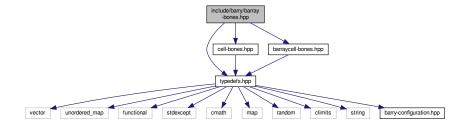
• include/barry/typedefs.hpp

Chapter 8

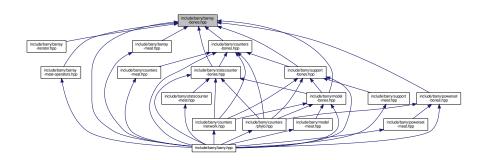
File Documentation

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

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



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



Classes

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

Macros

• #define BARRAY_BONES_HPP 1

8.1.1 Macro Definition Documentation

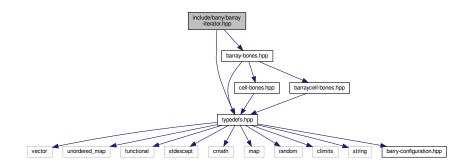
8.1.1.1 BARRAY_BONES_HPP

```
#define BARRAY_BONES_HPP 1
```

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

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

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



Classes

class ConstBArrayRowIter< Cell_Type, Data_Type >

Macros

• #define BARRAY_ITERATOR_HPP 1

8.2.1 Macro Definition Documentation

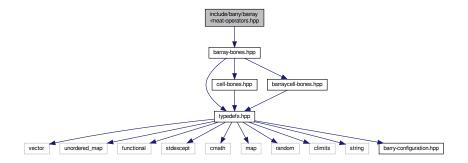
8.2.1.1 BARRAY_ITERATOR_HPP

#define BARRAY_ITERATOR_HPP 1

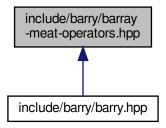
Definition at line 7 of file barray-iterator.hpp.

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 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 COL

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

8.3.1.2 ROW

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

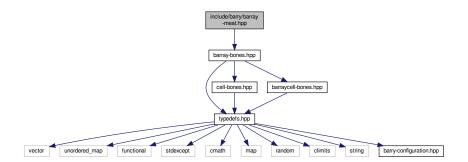
8.3.2 Function Documentation

8.3.2.1 checkdim_()

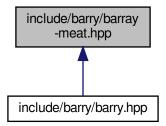
Definition at line 11 of file barray-meat-operators.hpp.

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

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

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

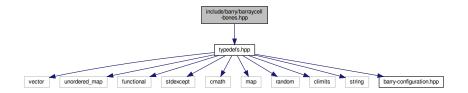
8.4.1.2 ROW

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

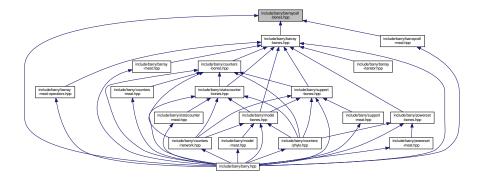
Definition at line 7 of file barray-meat.hpp.

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:

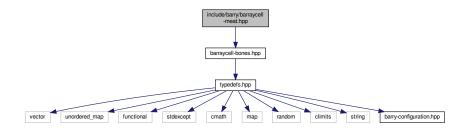


Classes

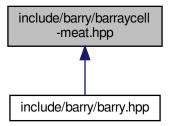
- class BArrayCell
 Cell_Type, Data_Type
- class BArrayCell_const< Cell_Type, Data_Type >

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

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

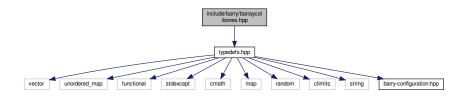


This graph shows which files directly or indirectly include this file:

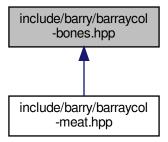


8.7 include/barry/barraycol-bones.hpp File Reference

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



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

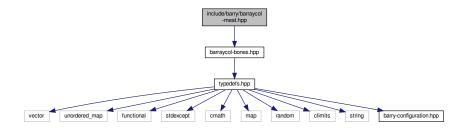


Classes

- class BArrayCol< Cell_Type, Data_Type >
- class BArrayCol_const< Cell_Type, Data_Type >

8.8 include/barry/barraycol-meat.hpp File Reference

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



Macros

• #define BARRY_BARRAYCOL_MEAT_HPP 1

8.8.1 Macro Definition Documentation

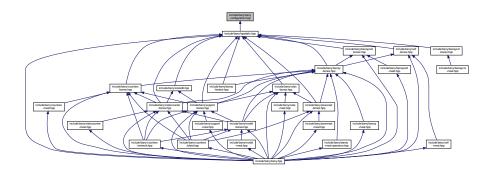
8.8.1.1 BARRY_BARRAYCOL_MEAT_HPP

#define BARRY_BARRAYCOL_MEAT_HPP 1

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

8.9 include/barry/barry-configuration.hpp File Reference

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



Macros

• #define BARRY_MAX_NUM_ELEMENTS static_cast< unsigned int >(UINT_MAX/2u)

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_CHECK_FINITE When specified, it will introduce a macro

```
• #define BARRY_SAFE_EXP -100.0
```

- #define BARRY_ISFINITE(a)
- #define BARRY_CHECK_SUPPORT(x, maxs)
- template<typename Ta , typename Tb >
 using Map = std::map< Ta, Tb >

8.9.1 Macro Definition Documentation

8.9.1.1 BARRY_CHECK_SUPPORT

```
#define BARRY_CHECK_SUPPORT(
          x,
          maxs )
```

Definition at line 46 of file barry-configuration.hpp.

8.9.1.2 BARRY ISFINITE

Definition at line 39 of file barry-configuration.hpp.

8.9.1.3 BARRY_MAX_NUM_ELEMENTS

```
\#define BARRY_MAX_NUM_ELEMENTS static_cast< unsigned int >(UINT_MAX/2u)
```

Definition at line 5 of file barry-configuration.hpp.

8.9.1.4 BARRY SAFE EXP

```
#define BARRY_SAFE_EXP -100.0
```

Definition at line 32 of file barry-configuration.hpp.

8.9.2 Typedef Documentation

8.9.2.1 Map

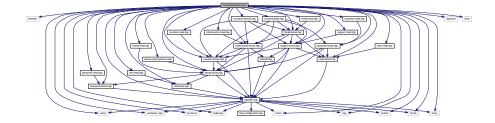
```
template<typename Ta , typename Tb >
using Map = std::map<Ta,Tb>
```

Definition at line 26 of file barry-configuration.hpp.

8.10 include/barry/barry.hpp File Reference

```
#include <iostream>
#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 "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 COUNTER_FUNCTION(a)
- #define COUNTER LAMBDA(a)
- #define RULE_FUNCTION(a)
- #define RULE_LAMBDA(a)

8.10.1 Macro Definition Documentation

8.10.1.1 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 66 of file barry.hpp.

8.10.1.2 COUNTER LAMBDA

Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
Counter_fun_type<Array_Type, Data_Type> a = \
[](const Array_Type & Array, uint i, uint j, Data_Type * data)
```

Definition at line 69 of file barry.hpp.

8.10.1.3 RULE FUNCTION

```
#define RULE_FUNCTION( a )
```

Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
inline bool (a) (const Array_Type & Array, uint i, uint j, Data_Type * data) \
```

Definition at line 73 of file barry.hpp.

8.10.1.4 RULE LAMBDA

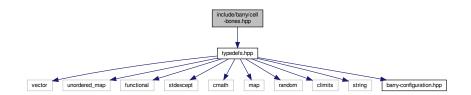
Value:

```
template <typename Array_Type = barry::BArray<>, typename Data_Type = bool> \
Rule_fun_type<Array_Type, Data_Type> a = \
[](const Array_Type & Array, uint i, uint j, Data_Type * data)
```

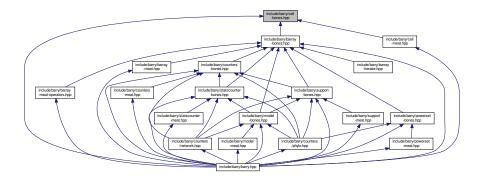
Definition at line 76 of file barry.hpp.

8.11 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:

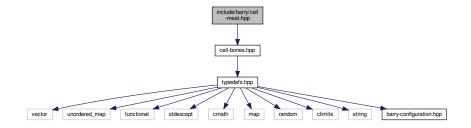


Classes

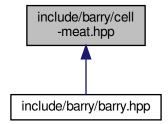
class Cell
 Cell_Type >
 Entries in BArray. For now, it only has two members:

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

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



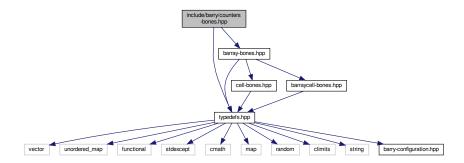
This graph shows which files directly or indirectly include this file:



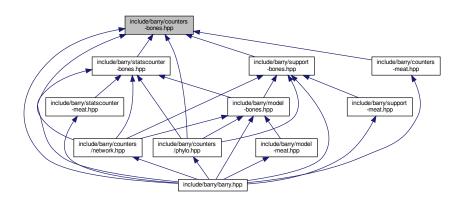
8.13 include/barry/col-bones.hpp File Reference

8.14 include/barry/counters-bones.hpp File Reference

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



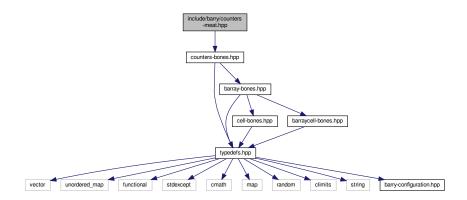
This graph shows which files directly or indirectly include this file:



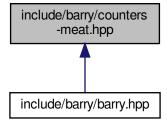
Classes

8.15 include/barry/counters-meat.hpp File Reference

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



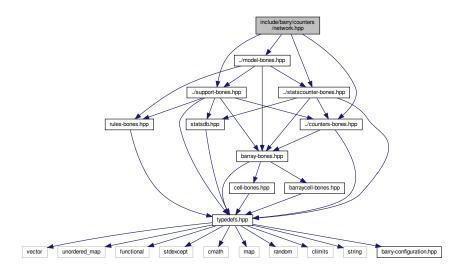
This graph shows which files directly or indirectly include this file:



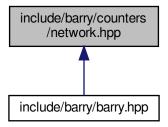
8.16 include/barry/counters/network.hpp File Reference

```
#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
 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.
```

8.16.1 Macro Definition Documentation

8.16.1.1 **NET_C_DATA_IDX**

Definition at line 79 of file network.hpp.

8.16.1.2 NET_C_DATA_NUM

Definition at line 80 of file network.hpp.

8.16.1.3 NETWORK_COUNTER

Value:

```
inline double (a) \
(const Network & Array, uint i, uint j, NetCounterData * data)
```

Function for definition of a network counter function

Definition at line 101 of file network.hpp.

8.16.1.4 NETWORK_COUNTER_LAMBDA

Lambda function for definition of a network counter function

Definition at line 104 of file network.hpp.

8.16.1.5 NETWORK_RULE

Function for definition of a network counter function

Definition at line 113 of file network.hpp.

8.16.1.6 NETWORK_RULE_LAMBDA

Lambda function for definition of a network counter function

Definition at line 116 of file network.hpp.

8.16.2 Typedef Documentation

8.16.2.1 NetCounter

```
typedef Counter<Network, NetCounterData > NetCounter
```

Definition at line 88 of file network.hpp.

8.16.2.2 NetCounters

typedef Counters< Network, NetCounterData> NetCounters

Definition at line 89 of file network.hpp.

8.16.2.3 NetModel

typedef Model<Network, NetCounterData> NetModel

Definition at line 92 of file network.hpp.

8.16.2.4 NetRule

typedef Rule<Network,bool> NetRule

Definition at line 93 of file network.hpp.

8.16.2.5 NetRules

typedef Rules<Network,bool> NetRules

Definition at line 94 of file network.hpp.

8.16.2.6 NetStatsCounter

typedef StatsCounter<Network, NetCounterData> NetStatsCounter

Definition at line 91 of file network.hpp.

8.16.2.7 NetSupport

typedef Support<Network, NetCounterData > NetSupport

Definition at line 90 of file network.hpp.

8.16.2.8 Network

```
typedef BArray<double, NetworkData> Network
```

Definition at line 87 of file network.hpp.

8.16.3 Function Documentation

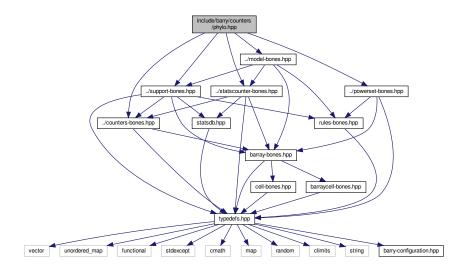
8.16.3.1 rules_zerodiag()

Number of edges.

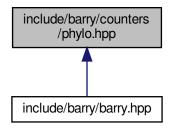
Definition at line 742 of file network.hpp.

8.17 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.

Macros

- #define PHYLO COUNTER(a)
 - Extension of a simple counter.
- #define PHYLO COUNTER LAMBDA(a)
- #define PHYLO_CHECK_MISSING()

Typedefs

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 Support < PhyloArray, PhyloCounterData, PhyloRuleData, uint > PhyloSupport
- typedef StatsCounter< PhyloArray, PhyloCounterData > PhyloStatsCounter
- typedef Model < PhyloArray, PhyloCounterData, PhyloRuleData, uint > 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.

- $\bullet \ \ void\ counter_gains\ (PhyloCounters\ *counters,\ std::vector < uint > nfun,\ bool\ duplication=true) \\$
 - 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_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.
- #define PHYLO_C_DATA_IDX(i) (data.operator[](i))
- typedef std::vector< uint > PhyloCounterData
- typedef std::vector< std::pair< uint, uint > > PhyloRuleData

8.17.1 Macro Definition Documentation

8.17.1.1 PHYLO_C_DATA_IDX

Definition at line 56 of file phylo.hpp.

8.17.1.2 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 data is nullptr.")
```

Definition at line 91 of file phylo.hpp.

8.17.1.3 PHYLO_COUNTER

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 85 of file phylo.hpp.

8.17.1.4 PHYLO_COUNTER_LAMBDA

Definition at line 88 of file phylo.hpp.

8.17.2 Typedef Documentation

8.17.2.1 PhyloArray

```
typedef BArray<uint, NodeData> PhyloArray
```

Definition at line 63 of file phylo.hpp.

8.17.2.2 PhyloCounter

```
typedef Counter<PhyloArray, PhyloCounterData > PhyloCounter
```

Definition at line 64 of file phylo.hpp.

8.17.2.3 PhyloCounterData

typedef std::vector< uint > PhyloCounterData

Definition at line 53 of file phylo.hpp.

8.17.2.4 PhyloCounters

typedef Counters< PhyloArray, PhyloCounterData> PhyloCounters

Definition at line 65 of file phylo.hpp.

8.17.2.5 PhyloModel

typedef Model<PhyloArray, PhyloCounterData, PhyloRuleData, uint > PhyloModel

Definition at line 70 of file phylo.hpp.

8.17.2.6 PhyloPowerSet

typedef PowerSet<PhyloArray, PhyloRuleData> PhyloPowerSet

Definition at line 71 of file phylo.hpp.

8.17.2.7 PhyloRule

typedef Rule<PhyloArray,PhyloRuleData> PhyloRule

Definition at line 66 of file phylo.hpp.

8.17.2.8 PhyloRuleData

typedef std::vector< std::pair< uint, uint > > PhyloRuleData

Definition at line 54 of file phylo.hpp.

8.17.2.9 PhyloRules

```
typedef Rules<PhyloArray,PhyloRuleData> PhyloRules
```

Definition at line 67 of file phylo.hpp.

8.17.2.10 PhyloStatsCounter

```
typedef StatsCounter<PhyloArray, PhyloCounterData> PhyloStatsCounter
```

Definition at line 69 of file phylo.hpp.

8.17.2.11 PhyloSupport

```
{\tt typedef~Support < PhyloArray,~PhyloCounterData,~PhyloRuleData,~uint > PhyloSupport}
```

Definition at line 68 of file phylo.hpp.

8.17.3 Function Documentation

8.17.3.1 get_last_name()

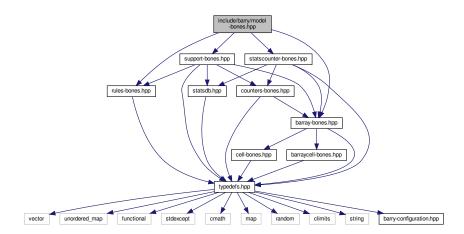
```
\begin{tabular}{ll} {\tt std::string get\_last\_name (} \\ {\tt bool } d\end{tabular} & [inline] \end{tabular}
```

Definition at line 96 of file phylo.hpp.

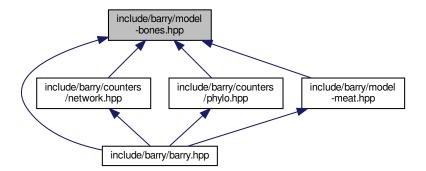
8.18 include/barry/model-bones.hpp File Reference

```
#include "barray-bones.hpp"
#include "support-bones.hpp"
#include "statscounter-bones.hpp"
#include "rules-bones.hpp"
```

Include dependency graph for model-bones.hpp:



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



Classes

class Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >

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

Functions

- 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)

```
    template<typename Array_Type >
        std::vector< double > keygen_default (const Array_Type &Array_)
```

Array Hasher class (used for computing support)

8.18.1 Function Documentation

8.18.1.1 keygen_default()

Array Hasher class (used for computing support)

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

8.18.1.2 likelihood_()

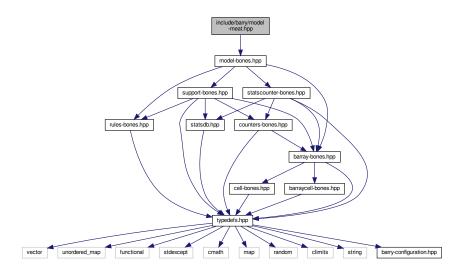
Definition at line 40 of file model-bones.hpp.

8.18.1.3 update_normalizing_constant()

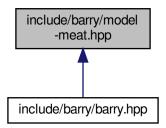
Definition at line 16 of file model-bones.hpp.

8.19 include/barry/model-meat.hpp File Reference

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



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



8.20 include/barry/models/geese.hpp File Reference

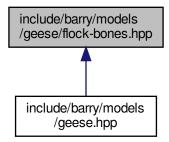
```
#include "geese/geese-node-bones.hpp"
#include "geese/geese-bones.hpp"
#include "geese/geese-meat.hpp"
#include "geese/geese-meat-constructors.hpp"
#include "geese/geese-meat-likelihood.hpp"
#include "geese/geese-meat-likelihood_exhaust.hpp"
#include "geese/geese-meat-simulate.hpp"
#include "geese/geese-meat-predict.hpp"
#include "geese/flock-bones.hpp"
```

#include "geese/flock-meet.hpp"
Include dependency graph for geese.hpp:



8.21 include/barry/models/geese/flock-bones.hpp File Reference

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



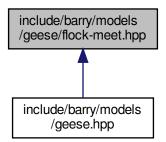
Classes

• class Flock

A Flock is a group of Geese.

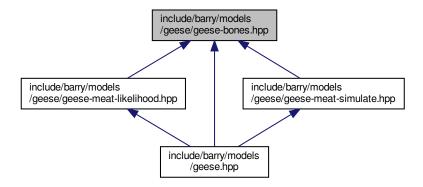
8.22 include/barry/models/geese/flock-meet.hpp File Reference

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



8.23 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)

8.23.1 Macro Definition Documentation

8.23.1.1 INITIALIZED

```
#define INITIALIZED( )

Value:
    if (!this->initialized) \
        throw std::logic_error("The model has not been initialized yet.");
```

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

8.23.2 Function Documentation

8.23.2.1 keygen_full()

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

8.23.2.2 RULE_FUNCTION()

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

8.23.2.3 vec_diff()

```
bool vec_diff (  {\it const std::} {\it vector} < {\it unsigned int} > \& s, \\ {\it const std::} {\it vector} < {\it unsigned int} > \& a ) \quad [inline]
```

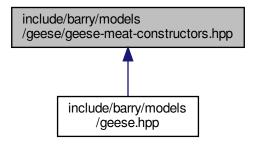
Definition at line 51 of file geese-bones.hpp.

8.23.2.4 vector_caster()

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

8.24 include/barry/models/geese/geese-meat-constructors.hpp File Reference

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



8.25 include/barry/models/geese/geese-meat-likelihood.hpp File Reference

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

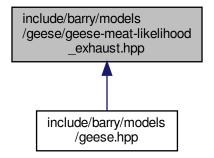


This graph shows which files directly or indirectly include this file:



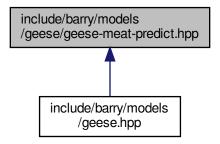
8.26 include/barry/models/geese/geese-meat-likelihood_exhaust.hpp File Reference

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



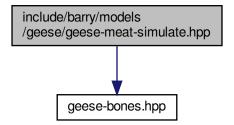
8.27 include/barry/models/geese/geese-meat-predict.hpp File Reference

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

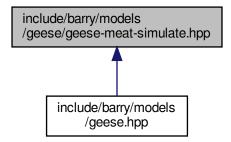


8.28 include/barry/models/geese/geese-meat-simulate.hpp File Reference

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

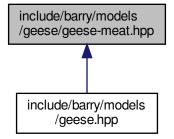


This graph shows which files directly or indirectly include this file:



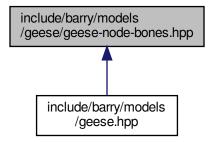
8.29 include/barry/models/geese/geese-meat.hpp File Reference

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



8.30 include/barry/models/geese/geese-node-bones.hpp File Reference

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



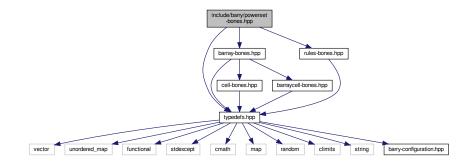
Classes

• class Node

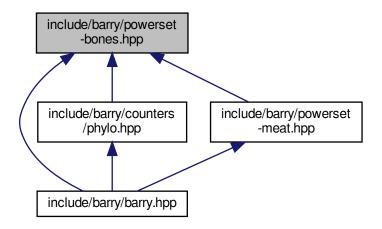
A single node for the model.

8.31 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:
```



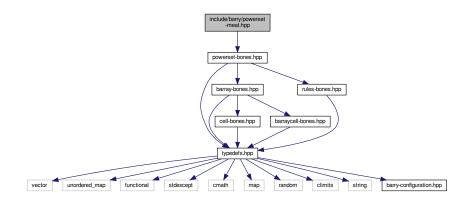
This graph shows which files directly or indirectly include this file:



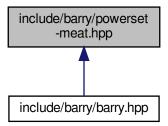
Classes

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

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

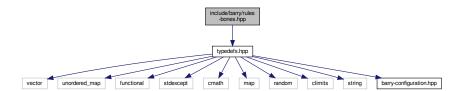


This graph shows which files directly or indirectly include this file:

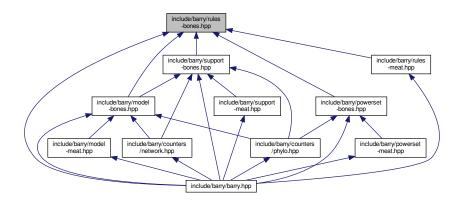


8.33 include/barry/rules-bones.hpp File Reference

#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)

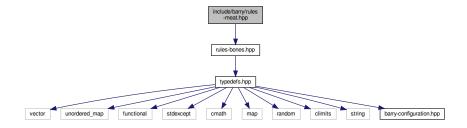
8.33.1 Function Documentation

8.33.1.1 rule_fun_default()

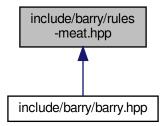
Definition at line 10 of file rules-bones.hpp.

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

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



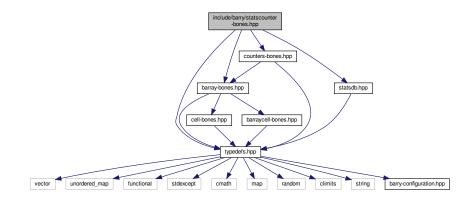
This graph shows which files directly or indirectly include this file:



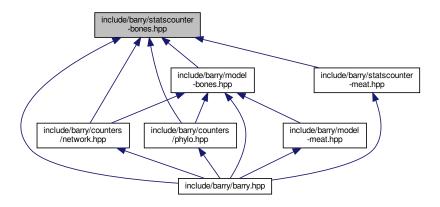
8.35 include/barry/statscounter-bones.hpp File Reference

```
#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:

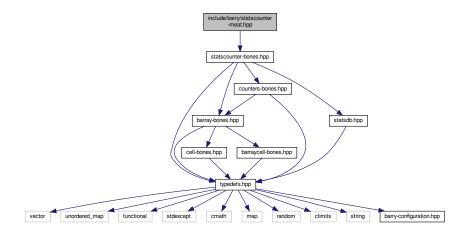


Classes

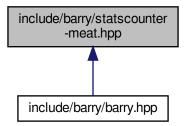
class StatsCounter < Array_Type, Data_Type >
 Count stats for a single Array.

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

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

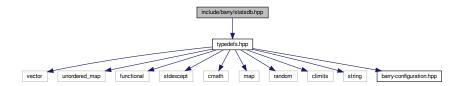


This graph shows which files directly or indirectly include this file:

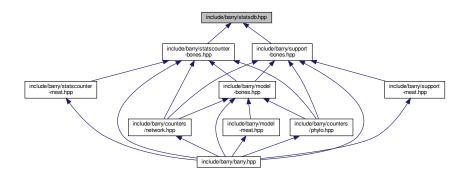


8.37 include/barry/statsdb.hpp File Reference

#include "typedefs.hpp"
Include dependency graph for statsdb.hpp:



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



Classes

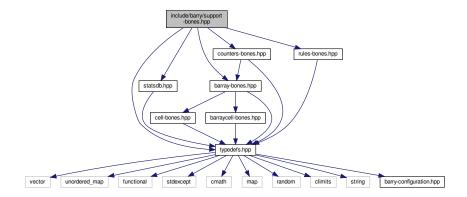
class FreqTable
 T >

Database of statistics.

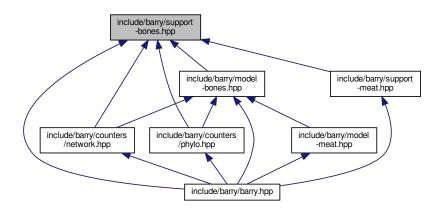
8.38 include/barry/support-bones.hpp File Reference

```
#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:

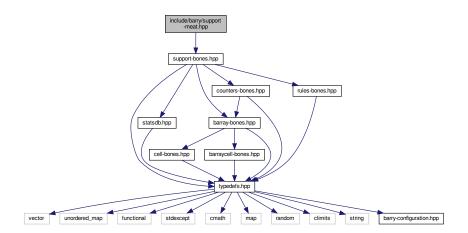


Classes

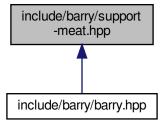
class Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >
 Compute the support of sufficient statistics.

8.39 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

8.39.1 Macro Definition Documentation

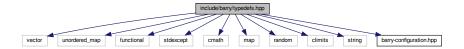
8.39.1.1 BARRY_SUPPORT_MEAT_HPP

#define BARRY_SUPPORT_MEAT_HPP 1

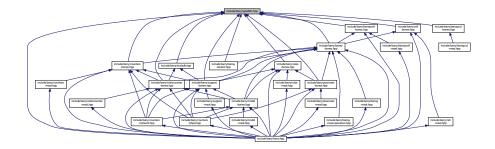
Definition at line 4 of file support-meat.hpp.

8.40 include/barry/typedefs.hpp File Reference

```
#include <vector>
#include <unordered_map>
#include <functional>
#include <stdexcept>
#include <cmath>
#include <map>
#include <random>
#include <climits>
#include <string>
#include "barry-configuration.hpp"
Include dependency graph for typedefs.hpp:
```



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



Classes

- class Entries < Cell_Type >
 - A wrapper class to store source, target, val from a BArray object.
- struct vecHasher< T >

Namespaces

CHECK

Integer constants used to specify which cell should be check.

• EXISTS

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

Typedefs

```
typedef unsigned int uint
typedef std::vector< std::pair< std::vector< double >, uint > > Counts_type
template<typename Cell_Type > using Row_type = Map< uint, Cell< Cell_Type > >
template<typename Cell_Type > using Col_type = Map< uint, Cell< Cell_Type > * >
template<typename Ta = double, typename Tb = uint> using MapVec_type = std::unordered_map< std::vector< Ta >, Tb, vecHasher< Ta > >
template<typename Array_Type , typename Data_Type > using Counter_fun_type = std::function< double(const Array_Type &, uint, uint, Data_Type *)> Counter and rule functions.
template<typename Array_Type , typename Data_Type >
```

using Rule_fun_type = std::function< bool(const Array_Type &, uint, uint, Data_Type *)>

Functions

```
    template < typename T >
        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::TWO = 1
const int EXISTS::UKNOWN = -1
const int EXISTS::AS_ZERO = 0
const int EXISTS::AS_ONE = 1
```

8.40.1 Typedef Documentation

8.40.1.1 Col_type

```
template<typename Cell_Type >
using Col_type = Map< uint, Cell<Cell_Type>* >
```

Definition at line 61 of file typedefs.hpp.

8.40.1.2 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 133 of file typedefs.hpp.

8.40.1.3 Counts_type

```
typedef std::vector< std::pair< std::vector<double>, uint > > Counts_type
```

Definition at line 54 of file typedefs.hpp.

8.40.1.4 MapVec_type

```
template<typename Ta = double, typename Tb = uint>
using MapVec_type = std::unordered_map< std::vector< Ta >, Tb, vecHasher<Ta> >
```

Definition at line 115 of file typedefs.hpp.

8.40.1.5 Row_type

```
template<typename Cell_Type >
using Row_type = Map< uint, Cell<Cell_Type> >
```

Definition at line 58 of file typedefs.hpp.

8.40.1.6 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 136 of file typedefs.hpp.

8.40.1.7 uint

```
typedef unsigned int uint
```

Definition at line 20 of file typedefs.hpp.

8.40.2 Function Documentation

8.40.2.1 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 147 of file typedefs.hpp.

8.40.2.2 vec_equal_approx()

Definition at line 165 of file typedefs.hpp.

8.40.2.3 vec_inner_prod()

Definition at line 185 of file typedefs.hpp.

8.41 README.md File Reference

Index

```
\simBArray
                                                               131
    BArray< Cell_Type, Data_Type >, 29
                                                     add
\simBArrayCell
                                                          Cell < Cell Type >, 52
    BArrayCell< Cell_Type, Data_Type >, 40
                                                          FreqTable < T >, 72
~BArrayCell const
                                                     add array
    BArrayCell_const< Cell_Type, Data_Type >, 42
                                                          Model<
                                                                      Array Type,
                                                                                      Data Counter Type,
\simBArrayCol
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
    BArrayCol < Cell_Type, Data_Type >, 45
                                                               87
~BArrayCol const
                                                     add_counter
    BArrayCol_const< Cell_Type, Data_Type >, 48
                                                          Counters < Array_Type, Data_Type >, 61, 62
\simCell
                                                                                      Data_Counter_Type,
                                                                      Array_Type,
    Cell< Cell Type >, 51
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
\simConstBArrayRowIter
                                                               87,88
    ConstBArrayRowlter < Cell_Type, Data_Type >, 55
                                                          StatsCounter < Array_Type, Data_Type >, 125
\simCounter
                                                          Support<
                                                                       Array Type,
                                                                                      Data Counter Type,
    Counter< Array Type, Data Type >, 58
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
\simCounters
    Counters < Array_Type, Data_Type >, 61
                                                      add data
\simEntries
                                                          Flock, 67
    Entries < Cell Type >, 64
                                                     add_rule
\simFlock
                                                                                      Data_Counter_Type,
                                                          Model<
                                                                      Array_Type,
    Flock, 66
                                                               Data Rule Type, Data Rule Dyn Type >,
\simFreqTable
                                                               88, 89
    FreqTable < T >, 71
                                                          PowerSet < Array_Type, Data_Rule_Type >, 113
\simGeese
                                                          Rules < Array Type, Data Type >, 121
    Geese, 76
                                                                      Array Type,
                                                                                      Data Counter Type,
                                                          Support<
\simModel
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
    Model <
                Array_Type,
                                Data Counter Type,
                                                               131, 132
         Data_Rule_Type, Data_Rule_Dyn_Type >,
                                                      add_rule_dyn
         87
                                                          Model <
                                                                      Array_Type,
                                                                                      Data Counter Type,
\simNetCounterData
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
    NetCounterData, 100
\simNetworkData
                                                          Support<
                                                                       Array Type,
                                                                                      Data Counter Type,
    NetworkData, 103
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
\simNode
    Node, 105
                                                      annotations
\simNodeData
                                                          Node, 106
    NodeData, 109
                                                     Array
\simPowerSet
                                                          ConstBArrayRowIter < Cell_Type, Data_Type >, 55
    PowerSet < Array_Type, Data_Rule_Type >, 113
                                                          StatsCounter< Array_Type, Data_Type >, 126
\simRule
                                                     array
    Rule < Array_Type, Data_Type >, 118
                                                          Node, 106
\simRules
                                                     array frequency
    Rules < Array_Type, Data_Type >, 120
                                                          Model<
                                                                      Array Type,
                                                                                      Data Counter Type,
\simStatsCounter
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
    StatsCounter< Array_Type, Data_Type >, 124
\simSupport
                 Array_Type,
                                                      arravs
                                Data_Counter_Type,
    Support<
                                                          Node, 106
         Data_Rule_Type, Data_Rule_Dyn_Type >,
                                                      arrays2support
```

Model< Array_Type, Data_Rule_Type, Data_Rule_Dyn_Type > 95 95 AS_ONE EXISTS, 23 as_vector FreqTable <t 7="">, 72 AS_ZERO EXISTS, 23 BArrayCell_Type, Data_Type >, 28, 29 BArrayCell_Type, Data_Type >, 25 ~BArrayCell_Type, Data_Type >, 25 ABArrayCell_Const< Cell_Type, Data_Type >, 38 BArrayCell_Const< Cell_Type, Data_Type >, 38 BArrayCell_Const< Cell_Type, Data_Type >, 38 Cetar, 29 col, 29 D, 30 default_val, 30 get_coll, 30</t>	Model Array Type Date Counter Type	harray-moat-operators has
Solution		• • • • • • • • • • • • • • • • • • • •
AS_ONE EXISTS, 23 as_vector FreqTable< T >, 72 AS_ZERO EXISTS, 23 BArray BArray< Cell_Type, Data_Type >, 28, 29 BArray BARRAY BONES BArray BARRAY BONES BARRAY BONES BARRAY BONES BARRAY BONES BARRAY BONES BARRAY BONES BARRAY BONES BARRAY BONES BARRAY BONES BARRAY BONES BARRAY BONES BARRAY BONES BARRAY BONES BARRAY BONES BARRAY BARRAY BONES BARRAY BARRAY BONES BARRAY BARRAY BONES BARRAY <b< td=""><td></td><td>_</td></b<>		_
EXISTS, 23 as_vector FreqTable< T >, 72 AS ZERO EXISTS, 23 BArray BArray Cell_Type, Data_Type >, 28, 29 BArray Cell_Type, Data_Type >, 25 ~BArray BArray (Cell_Type, Data_Type >, 28 BArray Cell_Cell_Type, Data_Type >, 28 BArray Cell_Cell_Type, Data_Type >, 38 BArray Cell_Const Cell_Type, Data_Type >, 42 BArray Cell_Const Cell_Type, Data_Type >, 43 BArray Cell_Const Cell_Type, Data_Type >, 44 BArray Cell_Const Cell_Type, Data_Type >, 44 BArray Cell_Const Cell_Type, Data_Type >, 45 BArray Cell_Const Cell_Type, Data_Type >, 45 BArray Cell_Const Cell_Type, Data_Type >, 46 BArray Cell_Const Cell_Type, Data_Type >, 47 BArray Cell_Const Cell_Type, Data_Type >, 48 BArray Cell_Const Cell_Type, Data_Type >, 49 BArray Cell_Const Cell_Type, Data_Type >, 49 BArray Cell_Const		
as vector FreqTable < T > , 72 AS ZERO EXISTS, 23 BArray BArray Cell_Type, Data_Type > , 28, 29 BArray Cell_Type, Data_Type > , 25	-	
Exists, 23 BArray BArray < Cell_Type, Data_Type >, 28, 29 BArray < Cell_Type, Data_Type >, 25	,	· · · · · · · · · · · · · · · · · · ·
AS_ZERO EXISTS, 23 BArray BArrayCell_Type, Data_Type >, 28, 29 BArrayCell_Type, Data_Type >, 28 BArrayCell_Type, Data_Type >, 28 BArrayCell_Constyc_Data_Type >, 38 BArrayCell_Const <cell_type, data_type="">, 38 BArrayCell_Const<cell_type, data_type="">, 38 clear, 29 D, 30 default_val, 30 get_coll, 30 get_co</cell_type,></cell_type,>		
EXISTS, 23 BArray BArray Cell_Type, Data_Type >, 28, 29 BArray Cell_Type, Data_Type >, 28 BArray Cell_Type, Data_Type >, 28 BArray Cell_Const-Cell_Type, Data_Type >, 38 BArrayCell_Const-Cell_Type, Data_Type >, 38 BArrayCell_Const-Cell_Type, Data_Type >, 38 Clear, 29 col, 29 D, 30 default val, 30 get_col, 30 get_col, 30 get_col, 30 get_col, 30 get_row, vec, 31 insert_cell, 32 is_empty, 32 non. 32 non. 32 non. 32 non. 33 operator*=, 34 operator*=, 34 operator*=, 35 out_of_range, 35 print, 35 reserve, 35 reserve, 35 resize, 36 rm_cell, 36 row, 36 set_data, 36 swap_colls, 37 swap_rows, 37 toggle_cold, 37 transpose, 38 barray-Dones.hpp BARRAY_BONES_HPP, 142 BArrayCol_const, 42 operator*=, 44 operator*=, 43 operator*=, 43 operator*=, 43 operator*=, 44 operator*=, 45 barrayCol_const. Cell_Type, Data_Type >, 48 BArrayCol_const. Cell_Type, Data_Type >, 48 BArrayCol_const. Cell_Type, 45 operator*=, 46 operator*=, 47 operator*=, 48 BArrayCol_const. Cell_Type, Data_Type >, 48 BArrayCol_const. Cell_Type, Data_Type >, 48 BArrayCol_const. Cell_Type, Data_Type >, 49 call Type, Data_Type >, 49 cal	•	
BArray Cell Type, Data Type >, 28, 29 BArray Cell Type, Data Type >, 25		
BArray Cell_Type, Data_Type >, 28, 29 BArray-Cell_Type, Data_Type >, 28, 29 BArray-Cell_Type, Data_Type >, 28 BArray-Cell_Type, Data_Type >, 38 BArray-Cell_Const-Cell_Type, Data_Type >, 38 BArray-Cell_Const-Cell_Type, Data_Type >, 38 Clear, 29 Col, 29 Co	2,40,10,20	• • • • • • • • • • • • • • • • • • • •
BArray Cell Type, Data Type >, 28, 29 BArray Cell Type, Data Type >, 25	BArray	
BArray< Cell_Type, Data_Type >, 25	BArray< Cell_Type, Data_Type >, 28, 29	
~BArray, 29 BArray, 28, 29 BArrayCell Cell_Type, Data_Type >, 38 BArrayCell_const < Cell_Type, Data_Type >, 38 BArrayCell_const < Cell_Type, Data_Type >, 38 BArrayCell_const < Cell_Type, Data_Type >, 38 BArrayCell, 30 get_col, 29 D, 30 default_val, 30 get_col, 30 get_col, 30 get_col, 30 get_col, 29 BArrayCell_const < Cell_Type, Data_Type >, 38 BArrayCell, 40 operator=, 40 operator=, 40 operator=, 41 operator=, 42 BArrayCell_const < Cell_Type, Data_Type >, 42 BArrayCol_const < Cell_Type, Data_Type >, 42 BArrayCol_const < Cell_Type, Data_Type >, 44 Deperator=, 43 Depe	BArray< Cell_Type, Data_Type >, 25	
BArray Cell Coll Type, Data_Type >, 38 BArrayCell_const< Cell_Type, Data_Type >, 38 clear, 29 col, 29 col, 29 col, 29 col, 30 default_val, 30 get_col, 30 get_col, 30 get_col, 30 get_col, 30 get_col, 30 get_col, 31 get_row, 31 get_row, 31 get_row, 31 get_row, 31 get_row, 32 nozero, 33 nrow, 33 operator =, 33 operator +, 33 operator -, 34 operator -, 35 out_of_range, 35 print, 35 reserve, 35 resize, 36 rm_cell, 36 row, 36 set_data, 36 swap_colls, 37 swap_rows, 37 toggle_lock, 37 transpose, 38 visited, 39 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-literator.hpp BARRAY_BONES_HPP, 142 barray-literator.hpp BARRAY_BONES_HPP, 142 barray-deleronst, 48 BArrayCol_const, 49 BARRAY_BONES_HPP, 142 barray-deleronst, 40 operator -=, 41 operator -=, 42 operator Cell_Type, Data_Type >, 42 BArrayCell_const, Cell_Type, Data_Type >, 42 BArrayCell_const, 42 operator <=, 43 BArrayCol, Const. Cell_Type, Data_Type >, 45 BArrayCol, 45 begin, 45 operator ==, 46 operator ==, 47 barraycol-meat.hpp BARRAY_BARRAYCOL_MEAT_HPP, 148 BArrayCol_const, 48 BArrayCol_const, 48 BArrayCol, 48 operator ==, 49 operator <=, 40 operator <=, 41 operator <=, 42 BArrayCell_const Cell_Type, Data_Type >, 42 ender <=, 43 operator	\sim BArray, 29	
BArrayCell< cell Type, Data Type >, 38 BArrayCell< const< Cell Type, Data Type >, 38 clear, 29 col, 29 col, 29 D, 30 default_val, 30 gel_coll, 31 gel_frow_vec, 31 insert_cell, 32 is_empty, 32 ncol, 32 nnozero, 33 nrow, 33 operator==, 33 operator==, 33 operator==, 33 operator==, 34 operator==, 34 operator==, 35 ou_of_arange, 35 print, 35 reserve, 36 reserve, 37 swap_colls, 37 swap_colls, 37 swap_colls, 37 resepose, 38 visited, 39 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-interator.hpp	BArray, 28, 29	
BArrayCell_const< Cell_Type, Data_Type > , 38 clear, 29 col, 29 D, 30 default_val, 30 get_col, 30 get_col, 30 get_col, 30 get_col, 30 get_col, 30 get_col, 30 get_row, 31 get_row, 21 get_row, 21 get_row, 22 is_empty, 32 ncol, 32 nnozero, 33 nrow, 33 operator+=, 33 operator+=, 33 operator+=, 33 operator+=, 33 operator+=, 33 operator-=, 34 operator-=, 34 operator-=, 34 operator-=, 35 operator-=, 36 operator-=, 36 operator-=, 37 swap_cols, 37 swap_cols, 37 swap_cols, 37 swap_rows, 37 toggle_cell, 37 toggle_cell, 37 toggle_colk, 37 transpose, 38 visited, 39 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-licerator.hpp Defator=-, 41 Doperator=-, 42 BARRAY_BONES_HPP, 142 Doperator, 43 Doperator, 43 Doperator, 43 Doperator, 43 Doperator, 43 Doperator, 43 Doperator, 4	BArrayCell< Cell_Type, Data_Type >, 38	-
clear, 29	BArrayCell_const< Cell_Type, Data_Type >, 38	
col, 29 D, 30 default_val, 30 get_cell, 30 get_cell, 30 get_col, 30 get_col, 30 get_col, 30 get_col, 30 get_row, 31 get_row, 20 insert_cell, 32 insert_cell, 32 incor, 33 nrow, 33 operator=, 34 operator=, 35 operator=, 36 operator=, 36 operator=, 37 swap_cols, 37 swap_cols, 37 swap_cols, 37 swap_rows, 37 toggle_cell, 37 toggle_cold, 38 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-lenatorhop		
D, 30 default_val, 30 get_ccell, 30 get_ccell, 30 get_col, 31 get_entries, 31 get_ow, 31 get_row, vec, 31 get_row, vec, 31 get_row, vec, 31 get_row, 32 nozero, 33 nrow, 33 operators=, 33 operators=, 33 operators=, 33 operators=, 33 operators=, 33 operators=, 34 operators=, 34 operators=, 35 out_of_range, 35 print, 35 reserve, 35 resize, 36 rm_cell, 36 row, 36 set_data, 36 swap_colls, 37 swap_rows, 37 toggle_cell, 37 toggle_lock, 37 transpose, 38 visited, 39 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-leared, 30 poreator=, 48 operator=, 46 operator=, 46 operator=, 46 operator=, 47 >barrayCol_const, 48 operator=, 46 operator=, 46 operator=, 46 operator=, 46 operator=, 46 operator=, 46 operator=, 47 >barray-bones.hpp BARRAY_BONES_HPP, 142 barray-learator,hpp	col, 29	
default_val, 30 get_coll, 31 get_trow, 32 get_trow, 33 get_trow, 34 get_trow, 35 get_trow, 36 get_trow, 36 get_trow, 37 get_trow, 38 get_trow, 36 get_trow, 36 get_trow, 36 get_trow, 36 get_trow, 36 get_trow, 36 get_trow, 37 get_trow, 37 get_trow, 37 get_trow, 37 get_trow, 38 get_trow, 37 get_trow, 38 get_trow, 37 get_trow, 38 get_trow, 38 get_trow, 38 get_trow, 36 get_trow, 36 get_trow, 37 get_trow, 37 get_trow, 38 get_trow, 39 g	D, 30	•
get_col, 30 get_col, 30 get_col, 30 get_col_vec, 30, 31 get_entries, 31 get_entries, 31 get_row, 32 get_row, 32 get_row, 32 get_row, 33 get_row, 34 get_row, 34 get_row, 35 get_row, 35 get_row, 36 get_row, 42 ge	- ·	•
get_col_vec, 30, 31 get_entries, 31 get_entries, 31 get_row_vec, 31 insert_cell, 32 is_empty, 32 nocl, 32 nnozero, 33 nrow, 33 operator*=, 33 operator*=, 33 operator*=, 33 operator*=, 33, 34 operator*=, 34 operator*=, 35 out_of_range, 35 print, 35 reserve, 35 reserve, 35 resize, 36 rm_cell, 36 set_data, 36 set_data, 36 set_data, 36 set_data, 36 swap_cells, 37 swap_rows, 37 toggle_cell, 37 toggle		•
get_entries, 31 get_row, 31 get_row, 21 get_row vec, 31 get_row vec, 42 get_row, 42 ge	-	operator=, 41
get_row, 31 get_row_vec, 31 get_row_vec, 31 insert_cell, 32 is_empty, 32 ncol, 32 nnozero, 33 nrow, 33 operator*=, 33 operator*=, 33 operator*=, 34 operator*=, 35 out_of_range, 35 print, 35 reserve, 35 reserve, 35 row, 36 set_data, 36 svap_cells, 37 swap_rows, 37 toggle_cell, 37 toggle_cell, 37 toggle_lock, 37 transpoose, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp BARRAY_BONES_HPP, 142 barray-cell_const< Cell_Type, Data_Type >, 42	·	operator==, 41
get_row_vec, 31 insert_cell, 32 is_empty, 32 ncol, 32 nnozero, 33 nrow, 33 operator*=, 33 operator(), 33 operator-=, 34 operator-=, 35 operator-=, 35 operator-=, 35 operator-=, 35 operator-=, 35 operator-=, 35 reserve, 35 resize, 36 rm_cell, 36 row, 36 set_data, 36 swap_cells, 37 swap_rows, 37 toggle_coll, 38 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp BARRAY_BONES_HPP, 142 barray-iterator.hqp BARRAY_BONES_HPP, 142 barray-Coll_const< Cell_Type, Data_Type >, 48 BARRAY_BONES_HPP, 1	-	BArrayCell_const
insert_cell, 32 is_empty, 32 nnol, 32 nnozero, 33 nrow, 33 operator*=, 33 operator(), 33 operator*=, 34 operator*=, 35 operator*=, 35 out_of_range, 35 print, 35 reserve, 36 rm_cell, 36 row, 36 set_data, 36 swap_colls, 37 swap_rows, 37 toggle_cell, 37 toggle_cell, 37 toggle_lock, 37 transpose, 38 barray_bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp AARAY_BONES_HPP, 142 barray-iterator.hpp AARATAY_Coll_const, 42 barray-Coll_const, 42 barray		BArrayCell_const< Cell_Type, Data_Type >, 42
is_empty, 32 ncol, 32 nnozero, 33 nrow, 33 operator*=, 33 operator(), 33 operator-=, 34 operator-=, 35 out_of_range, 35 print, 35 reserve, 35 reserve, 35 reserve, 35 reserve, 36 rm_cell, 36 row, 36 set_data, 36 swap_cells, 37 swap_rows, 37 toggle_cell, 37 swap_rows, 37 toggle_cell, 37 toggle_lock, 37 transpose, 38 barrayCol_const< Cell_Type, Data_Type >, 48 barrayCol, 45 serizor, 46 operator-=, 47 barrayCol_const< Cell_Type, Data_Type >, 47 barrayCol-meat.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp		BArrayCell_const< Cell_Type, Data_Type >, 42
ncol, 32 nnozero, 33 nrow, 33 operator*=, 33 operator, 34 operator, 34 operator, 34 operator=, 34 operator=, 35 operator=, 35 operator=, 35 out_of_range, 35 print, 35 resize, 36 rm_cell, 36 row, 36 set_data, 36 swap_cells, 37 swap_rows, 37 toggle_lock, 37 transpose, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp BARRAY_BONES_HPP, 142 barray-col_cents, 43 operator, 44 operator, 43 operator, 45 BArrayCol, Cell_Type, Data_Type, 45 operator, 45 operator, 46 operator, 47 operator, 48 operator, 43		\sim BArrayCell_const, 42
nnozero, 33 nrow, 33 nrow, 33 operator*=, 33 operator(), 33 operator+=, 33, 34 operator-=, 34 operator-=, 34 operator-=, 35 operator-=, 35 out_of_range, 35 print, 35 reserve, 35 resize, 36 rm_cell, 36 row, 36 set_data, 36 set_data, 36 set_data, 36 swap_rows, 37 swap_rows, 37 toggle_cell, 37 toggle_cell, 37 toggle_lock, 37 transpose, 38 visited, 39 zero_col, 38 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-col, 48 Derator-=, 43 operator-=, 43 operator-=, 44 operator-=, 43 BArrayCol Cell_Type, Data_Type >, 45 BArrayCol RATrayCol Cell_Type, Data_Type >, 48 BArrayCol, 45 begin, 45 end, 45 operator C=, 46 operator-=, 47 barraycol-meat.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp		BArray< Cell_Type, Data_Type >, 38
operator*=, 33 operator*=, 33 operator*=, 33 operator*=, 33, 34 operator*=, 34 operator*=, 35 operator*=, 35 out_of_range, 35 print, 35 resize, 36 rm_cell, 36 set_data, 36 set_data, 36 set_data, 36 swap_colls, 37 swap_rows, 37 toggle_cell, 37 toggle_lock, 37 transpose, 38 visited, 39 zero_col, 38 zero_row, 38 barray Olegaror* barray operator*=, 43 operator*=, 43 operator*>=, 44 operator*=, 43 operator*>=, 45 operator*<=, 42 operator*Ocell_Type, Data_Type >, 48 operator*<=, 46 operator*=, 47 operator*=, 48 operator*=		BArrayCell_const, 42
operator*=, 33 operator(), 33 operator(), 33 operator-=, 34 operator-=, 34 operator-=, 34 operator-=, 35 operator-=, 35 out_of_range, 35 print, 35 reserve, 35 resize, 36 rm_cell, 36 row, 36 set_data, 36 swap_cells, 37 swap_rows, 37 toggle_cell, 37 toggle_cell, 37 toggle_lock, 37 transpose, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp Samaray and soperator (=, 48) operator (=, 48) operator (=, 43) operator (=, 43) operator (=, 43) operator (=, 43) operator (=, 44) operator (=, 44) operator (=, 45) sharrayCol (Cell_Type, Data_Type >, 45 BArrayCol_const (Cell_Type, Data_Type >, 48 BArrayCol, 45 BArrayCol, 45 BArrayCol, 45 begin, 45 operator Cell_Type, 45 operator (=, 46 operator-=, 46 op		operator Cell_Type, 43
operator(), 33 operator+=, 33, 34 operator-=, 34 operator-=, 34 operator-=, 35 operator==, 35 operator==, 35 out_of_range, 35 print, 35 reserve, 35 resize, 36 rm_cell, 36 row, 36 set_data, 36 swap_cells, 37 swap_rows, 37 toggle_cell, 37 toggle_cell, 37 toggle_cell, 37 toggle_col, 39 zero_col, 38 zero_row, 38 barrayCol SARRAY_BONES_HPP, 142 barray-iterator.hpp operator<=, 43 operator>=, 44 operator>=, 44 operator>=, 44 operator>=, 44 operator>=, 44 operator>=, 45 operator(cell_Type, Data_Type >, 45 BArrayCol SARrayCol BArrayCol BArrayCol BArrayCol, 45 BArrayCol, 45 begin, 45 operator Cell_Type, 45 operator Cell_Type, 45 operator>=, 46 operator>=, 46 operator>=, 46 operator>=, 46 operator==, 47 barray-olorest.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp operator<=, 48 operator<=, 48 BArrayCol_const		operator!=, 43
operator+=, 33, 34 operator-=, 34 operator/=, 34 operator/=, 35 operator/=, 35 operator/=, 35 out_of_range, 35 print, 35 reserve, 35 resize, 36 rm_cell, 36 row, 36 set_data, 36 swap_cells, 37 swap_rows, 37 toggle_cell, 37 toggle_lock, 37 transpose, 38 visited, 39 zero_row, 38 barray-iterator.hpp SARRAY_BONES_HPP, 142 operator>=, 44 operator>=, 44 operator>=, 44 operator>=, 45 operator>=, 46 operator==, 47 operator==, 46 operator==, 47 operator==, 46 operator==, 47 operator==, 46 operator==, 47 operator==, 46 operator==, 47 operator==, 47 operator==, 46 operator==, 47 operator==, 46 operator==, 47 operator==, 46 operator==, 47 operator==, 46 operator==, 47 operator==, 48 operator==, 49 operator==	•	•
operator-=, 34 operator/=, 34 operator/=, 34 operator/=, 35 operator==, 35 out_of_range, 35 print, 35 reserve, 35 resize, 36 rm_cell, 36 row, 36 set_data, 36 swap_cells, 37 swap_rows, 37 toggle_cell, 37 toggle_cell, 37 toggle_lock, 37 transpose, 38 visited, 39 zero_col, 38 zero_row, 38 barrayCol SARRAY_BONES_HPP, 142 barray-iterator.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp BARRAY_BONES_HPP, 142 barrayCol BArrayCol BArrayCol BArrayCol Cell_Type, Data_Type >, 48 BArrayCol, 45 Degrator+=, 46 Operator+=, 46 Operator+=, 46 Operator-=, 46 Operator-=, 46 Operator-=, 46 Operator-=, 47 barray-col_meat.hpp BARRAY_BONES_HPP, 142 BArrayCol_const, 48 BArrayCol, 48 Operator!=, 48	•	•
operator/=, 34 operator=, 35 operator==, 35 operator==, 35 out_of_range, 35 print, 35 reserve, 35 resize, 36 rm_cell, 36 row, 36 set_data, 36 swap_cells, 37 swap_rows, 37 toggle_cell, 37 toggle_lock, 37 transpose, 38 visited, 39 zero_col, 38 zero_row, 38 barrayCol BArrayCol BArrayCol BArrayCol BArrayCol BArrayCol Cell_Type, Data_Type >, 48 BArrayCol, 45 operator=, 46 operator=, 46 operator-=, 46 operator-=, 46 operator-=, 46 operator-=, 46 operator==, 47 barraycol-meat.hpp BARRAY_BONES_HPP, 142 BArrayCol_const< Cell_Type, Data_Type >, 47 ~BArrayCol_const< Cell_Type, Data_Type >, 47	•	•
operator=, 35 operator==, 35 operator==, 35 out_of_range, 35 print, 35 print, 35 reserve, 35 resize, 36 rm_cell, 36 row, 36 set_data, 36 swap_cells, 37 swap_rows, 37 toggle_cell, 37 toggle_lock, 37 transpose, 38 visited, 39 zero_col, 38 zero_row, 38 barrayCol server, 35 BarrayCol < Cell_Type, Data_Type >, 48 BArrayCol, 45 BArrayCol, 45 BArrayCol, 45 BArrayCol, 45 begin, 45 operator Cell_Type, 45 operator*=, 46 operator*=, 47 barraycol-meat.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp BARRAY_BARAY_Col_const, 48 operator=, 48	•	operator>=, 44
operator==, 35 out_of_range, 35 print, 35 print, 35 reserve, 35 resize, 36 rm_cell, 36 row, 36 set_data, 36 swap_cells, 37 swap_rows, 37 toggle_cell, 37 toggle_lock, 37 transpose, 38 visited, 39 zero_col, 38 zero_row, 38 barrayCol< Cell_Type, Data_Type >, 48 BArrayCol, 45 BArrayCol, 45 BArrayCol, 45 begin, 45 end, 45 operator Cell_Type, 45 operator Cell_Type, 45 operator*=, 46 operator*=, 47 barrayCol, 39 BARRAY_BONES_HPP, 142 barray-iterator.hpp	•	,
out_of_range, 35 print, 35 print, 35 print, 35 print, 35 preserve, 35 preserve, 36 prow, 37 prow, 38 prow, 38 prow, 39 prow, 38 prow, 39 prow, 38 prow, 30 prow, 45 prow, 30 prow, 30 prow, 45 p	·	-
print, 35 reserve, 35 resize, 36 rm_cell, 36 row, 36 set_data, 36 swap_cells, 37 swap_rows, 37 toggle_cell, 37 toggle_lock, 37 transpose, 38 visited, 39 zero_col, 38 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp BArrayCol< Cell_Type, Data_Type >, 44 ~BArrayCol, 45 BArrayCol, 45 BArrayCol, 45 begin, 45 end, 45 operator Cell_Type, 45 operator*=, 46 operator*=, 47 barray-col_const< Cell_Type, Data_Type >, 47 ~BArrayCol_const, 48 BArrayCol, 48 operator!=, 48	•	
reserve, 35 resize, 36 resize, 36 rm_cell, 36 row, 36 set_data, 36 set_data, 36 swap_cells, 37 swap_cols, 37 swap_rows, 37 toggle_cell, 37 toggle_lock, 37 transpose, 38 visited, 39 zero_col, 38 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp SARRAY_BARRAY_Col_MEAT_HPP, 148 SARRAY_BARRAY_Col_das SARRAY_Col_das SARR		
resize, 36 rm_cell, 36 row, 36 set_data, 36 swap_cells, 37 swap_cols, 37 swap_rows, 37 toggle_cell, 37 toggle_lock, 37 transpose, 38 visited, 39 zero_col, 38 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp BARRAY_BONES_HPP, 142 BARRAY_Col_const, 48 BARRAY_Col_d, 48 operator!=, 48	·	
rm_cell, 36 row, 36 set_data, 36 set_data, 36 swap_cells, 37 swap_rows, 37 swap_rows, 37 toggle_cell, 37 transpose, 38 visited, 39 zero_col, 38 zero_row, 38 begin, 45 end, 45 operator Cell_Type, 45 operator+=, 46 operator-=, 46 operator-=, 46 operator-=, 46 operator-=, 46 operator==, 47 barraycol-meat.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp begin, 45 operator **- operator		
row, 36 set_data, 36 set_data, 36 swap_cells, 37 swap_cols, 37 swap_rows, 37 toggle_cell, 37 toggle_lock, 37 transpose, 38 visited, 39 zero_col, 38 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp end, 45 operator Cell_Type, 45 operator*=, 46 operator*=, 46 operator*=, 46 operator*=, 46 operator*=, 46 operator*=, 47 barraycol-meat.hpp BARRAYCOL_MEAT_HPP, 148 BArrayCol_const< Cell_Type, Data_Type >, 47 ~BArrayCol_const, 48 BArrayCol, 48 operator!=, 48		•
set_data, 36 swap_cells, 37 swap_cols, 37 swap_rows, 37 toggle_cell, 37 toggle_lock, 37 visited, 39 zero_col, 38 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp swap_cols, 37 operator*=, 46 operator-=, 46 operator/=, 46 operator==, 47 operator-=, 46 operator-=, 46 operator-=, 47 operator-=, 46 operator-=, 46 operator-=, 46 operator-=, 47 operator-=, 48 ABARAY_BARRAYCOL_MEAT_HPP, 148 BARRAYCOl_const < Cell_Type, Data_Type >, 47 ~BArrayCol_const, 48 BArrayCol, 48 operator!=, 48		_
swap_cells, 37 swap_cols, 37 swap_rows, 37 toggle_cell, 37 toggle_lock, 37 transpose, 38 visited, 39 zero_col, 38 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp swap_cells, 37 operator*=, 46 operator-=, 46 operator-=, 46 operator-=, 46 operator-=, 47 operator-=, 46 operator-=, 47 operator-=, 48 operator-=, 48 operator-=, 48 operator-=, 48		
swap_cols, 37 swap_rows, 37 toggle_cell, 37 toggle_lock, 37 transpose, 38 visited, 39 zero_col, 38 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp swap_rows, 37 operator+=, 46 operator-=, 46 operator==, 47 operator-=, 46 operator-=, 48 operator-=, 48		
swap_rows, 37 toggle_cell, 37 toggle_lock, 37 transpose, 38 visited, 39 zero_col, 38 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp swap_rows, 37 operator=, 46 operator=, 46 operator=, 47 barraycol-meat.hpp BARRY_BARRAYCOL_MEAT_HPP, 148 BArrayCol_const< Cell_Type, Data_Type >, 47 ~BArrayCol_const, 48 BArrayCol, 48 operator!=, 48	•—	•
toggle_cell, 37 toggle_lock, 37 transpose, 38 visited, 39 zero_col, 38 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp speciator =, 46 operator=, 46 operator==, 47 barraycol-meat.hpp BARRY_BARRAYCOL_MEAT_HPP, 148 BArrayCol_const < Cell_Type, Data_Type >, 47 ~BArrayCol_const, 48 BArrayCol, 48 operator!=, 48	• —	•
toggle_lock, 37 transpose, 38 visited, 39 zero_col, 38 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp sportator=, 46 operator==, 47 barraycol-meat.hpp BARRAYCOL_MEAT_HPP, 148 BArrayCol_const < Cell_Type, Data_Type >, 47 ~BArrayCol_const, 48 BArrayCol, 48 operator!=, 48	• —	•
transpose, 38 visited, 39 zero_col, 38 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp sportator==, 47 barraycol-meat.hpp BARRAYCOL_MEAT_HPP, 148 BArrayCol_const< Cell_Type, Data_Type >, 47 ~BArrayCol_const, 48 BArrayCol, 48 operator!=, 48		•
visited, 39 zero_col, 38 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp barray-iterator.hpp barray-iterator.hpp barray-col_meat.hpp BARRAY_BARRAYCOL_MEAT_HPP, 148 BARRAY_BARRAYCOL_onst< Cell_Type, Data_Type >, 47 ~BArrayCol_const, 48 BArrayCol, 48 operator!=, 48	* * —	•
zero_col, 38 zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp BARRAY_BONES_HPP, 142	•	•
zero_row, 38 barray-bones.hpp BARRAY_BONES_HPP, 142 barray-iterator.hpp BARRAY_BONES_HPP, 142 BARRAY_BONES_HPP, 142 BARRAY_BONES_HPP, 142 BARRAY_BONES_HPP, 142 BARRAY_BONES_HPP, 142 BARRAY_BONES_HPP, 142		· · · · · · · · · · · · · · · · · · ·
barray-bones.hpp ~BArrayCol_const, 48 BARRAY_BONES_HPP, 142 BArrayCol, 48 barray-iterator.hpp operator!=, 48		
BARRAY_BONES_HPP, 142 barray-iterator.hpp BArrayCol, 48 operator!=, 48		
barray-iterator.hpp operator!=, 48	BARRAY_BONES_HPP, 142	· —
BARRAY_ITERATOR_HPP, 142	barray-iterator.hpp	
	BARRAY_ITERATOR_HPP, 142	•

operator<, 48	visited, 53
operator<=, 48	change_stats
operator>, 49	Support< Array_Type, Data_Counter_Type,
operator>=, 49	Data_Rule_Type, Data_Rule_Dyn_Type >,
operator==, 49	135
barry, 21	CHECK, 22
barry-configuration.hpp	BOTH, 22
BARRY_CHECK_SUPPORT, 149	NONE, 22
BARRY ISFINITE, 149	ONE, 22
BARRY_MAX_NUM_ELEMENTS, 149	TWO, 22
BARRY_SAFE_EXP, 150	checkdim_
Map, 150	barray-meat-operators.hpp, 144
barry.hpp	clear
COUNTER_FUNCTION, 151	BArray< Cell_Type, Data_Type >, 29
COUNTER_LAMBDA, 151	Counters < Array_Type, Data_Type >, 62
RULE_FUNCTION, 152	FreqTable $<$ T $>$, 72
RULE_LAMBDA, 152	Rules < Array_Type, Data_Type >, 121
barry::counters, 21	COL
barry::counters::network, 22	barray-meat-operators.hpp, 144
barry::counters::phylo, 22	barray-meat.hpp, 145
BARRY_BARRAYCOL_MEAT_HPP	col
barraycol-meat.hpp, 148	BArray< Cell_Type, Data_Type >, 29
BARRY_CHECK_SUPPORT	Col_type
barry-configuration.hpp, 149	typedefs.hpp, 187
BARRY_ISFINITE	ConstBArrayRowlter
barry-configuration.hpp, 149	ConstBArrayRowlter< Cell_Type, Data_Type >, 55
BARRY_MAX_NUM_ELEMENTS	ConstBArrayRowlter< Cell_Type, Data_Type >, 54
barry-configuration.hpp, 149	~ConstBArrayRowlter, 55
BARRY_SAFE_EXP	Array, 55
barry-configuration.hpp, 150	ConstBArrayRowlter, 55
BARRY_SUPPORT_MEAT_HPP	current_col, 55
support-meat.hpp, 185	current_row, 55
begin	iter, 56
BArrayCol< Cell_Type, Data_Type >, 45	coordinates_free
PowerSet< Array_Type, Data_Rule_Type >, 113	PowerSet < Array_Type, Data_Rule_Type >, 115
blengths	Support< Array_Type, Data_Titule_Type >, TTO
NodeData, 110	Data_Rule_Type, Data_Rule_Dyn_Type >,
BOTH	135
CHECK, 22	coordinates_locked
EXISTS, 23	PowerSet < Array_Type, Data_Rule_Type >, 115
calc	Support< Array_Type, Data_Counter_Type,
PowerSet< Array_Type, Data_Rule_Type >, 114	Data_Rule_Type, Data_Rule_Dyn_Type >,
Support< Array_Type, Data_Counter_Type,	135
Data_Rule_Type, Data_Rule_Dyn_Type >,	count
132	Counter< Array_Type, Data_Type >, 58
calc_reduced_sequence	count_all
·	StatsCounter< Array_Type, Data_Type >, 125
Geese, 76	count_current
calc_sequence	StatsCounter< Array_Type, Data_Type >, 125
Geese, 76	count_fun
Cell	Counter< Array_Type, Data_Type >, 59
Cell< Cell_Type >, 50–52	count_init
Cell< Cell_Type >, 49	StatsCounter< Array_Type, Data_Type >, 125
∼Cell, 51	Counter
add, 52	Counter< Array_Type, Data_Type >, 57
Cell, 50–52	Counter< Array_Type, Data_Type >, 56
operator Cell_Type, 53	~Counter, 58
operator=, 53	count, 58
value 53	,

count_fun, 59	Network counters, 13
Counter, 57	counter_neofun
data, 59	Phylo counters, 18
delete_data, 59	counter_neofun_a2b
desc, 59	Phylo counters, 18
init, 58	counter_nodecov
init_fun, 59	Network counters, 13
name, 59	counter_nodeicov
operator=, 58	Network counters, 13
counter_absdiff	counter_nodematch
Network counters, 11	Network counters, 13
counter_co_opt	counter_nodeocov
Phylo counters, 16	Network counters, 13
counter_cogain	counter_odegree
Phylo counters, 16	Network counters, 14
counter_ctriads	counter_odegree15
Network counters, 11	Network counters, 14
counter_degree	counter_ostar2
Network counters, 11	Network counters, 14
counter_deleted	counter_overall_changes
StatsCounter< Array_Type, Data_Type >, 126	Phylo counters, 18
counter_density	counter_overall_gains
Network counters, 11	Phylo counters, 19
counter_diff	counter_overall_loss
Network counters, 11	Phylo counters, 19
counter_edges	counter_subfun
Network counters, 12	Phylo counters, 19
counter_fun	counter_ttriads
Model	Network counters, 14
Data_Rule_Type, Data_Rule_Dyn_Type >,	Counters
95	Counters < Array_Type, Data_Type >, 61
Counter_fun_type	counters
typedefs.hpp, 188	Geese, 80
COUNTER_FUNCTION	Model < Array_Type, Data_Counter_Type
barry.hpp, 151	Data_Rule_Type, Data_Rule_Dyn_Type >
counter_gains	95
Phylo counters, 16	StatsCounter< Array_Type, Data_Type >, 127
counter_gains_k_offspring	Support< Array_Type, Data_Counter_Type
Phylo counters, 17	Data_Rule_Type, Data_Rule_Dyn_Type $>$
counter_genes_changing	135
Phylo counters, 17	Counters< Array_Type, Data_Type >, 60
counter_idegree	\sim Counters, 61
Network counters, 12	add_counter, 61, 62
counter_idegree15	clear, 62
Network counters, 12	Counters, 61
counter_isolates	operator=, 62
Network counters, 12	operator[], 62
counter_istar2	size, 63
Network counters, 12	counters_ptr
COUNTER_LAMBDA	Flock, 67
barry.hpp, 151	Counting, 9
counter_longest	Counts_type
Phylo counters, 17	typedefs.hpp, 188
counter_loss	current_col
Phylo counters, 17	ConstBArrayRowlter< Cell_Type, Data_Type >, 55
counter_maxfuns	current_row
Phylo counters, 18	ConstBArrayRowlter< Cell_Type, Data_Type >, 55
	Ourside Array rowner Com Type, Data Type 2, 3

StatsCounter< Array_Type, Data_Type >, 127 Support< Array_Type, Data_Counter_Type,	Entries < Cell_Type >, 63 ~Entries, 64 Entries, 64 resize, 64 source, 65 target, 65 val, 65 EXISTS, 23 AS_ONE, 23 AS_ZERO, 23 BOTH, 23
Counter< Array_Type, Data_Type >, 59 PowerSet< Array_Type, Data_Rule_Type >, 116 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 136	NONE, 24 ONE, 24 TWO, 24 UKNOWN, 24
default_val BArray< Cell_Type, Data_Type >, 30 delete_counters Geese, 80	first_calc_done Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 96
Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 136 delete_data	Flock, 65 ~Flock, 66 add_data, 67 counters_ptr, 67
Counter< Array_Type, Data_Type >, 59 delete_rengine Geese, 80	dat, 70 Flock, 66 init, 67
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 96 delete_rules	initialized, 70 likelihood_joint, 67 nfunctions, 70 nfuns, 68
Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 136 delete rules dyn	nleafs, 68 nnodes, 68 nterms, 68
Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 137	ntrees, 69 operator(), 69 rengine, 70 set_seed, 69
<pre>delete_support Geese, 81 desc Counter< Array_Type, Data_Type >, 59</pre>	support, 70 FreqTable FreqTable $T >$, 71 FreqTable $T >$, 71
directed NetworkData, 103 duplication	~FreqTable, 71 add, 72 as_vector, 72
Node, 107 NodeData, 110 EmptyArray	clear, 72 FreqTable, 71 get_data, 72 get_data_ptr, 72
PowerSet < Array_Type, Data_Rule_Type >, 116 StatsCounter < Array_Type, Data_Type >, 127 Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,	print, 72 reserve, 73 size, 73
137	Geese, 73
end BArrayCol< Cell_Type, Data_Type >, 45	~Geese, 76 calc_reduced_sequence, 76
PowerSet < Array_Type, Data_Rule_Type >, 114	calc_reduced_sequence, 76 calc_sequence, 76
Entries	counters, 80
Entries< Cell_Type >, 64	delete_counters, 80

delete_rengine, 80	FreqTable < T >, 72
delete_support, 81	PowerSet< Array_Type, Data_Rule_Type >, 114
Geese, 75, 76	get entries
get_probabilities, 76	BArray< Cell_Type, Data_Type >, 31
inherit support, 77	get_last_name
init, 77	phylo.hpp, 166
init_node, 77	get_norm_const
initialized, 81	Model< Array_Type, Data_Counter_Type,
likelihood, 77	Data_Rule_Type, Data_Rule_Dyn_Type >,
likelihood_exhaust, 77	90
map_to_nodes, 81	get parent
nfunctions, 81	Node, 106
nfuns, 78	get_probabilities
nleafs, 78	Geese, 76
nnodes, 78	get_pset
nodes, 81	Model Array_Type, Data_Counter_Type,
nterms, 78	Data Rule Type, Data Rule Dyn Type >,
observed_counts, 78	90
operator=, 78, 79	
predict, 79	get_row BArray< Cell_Type, Data_Type >, 31
predict_backend, 79	
print_observed_counts, 79	get_row_vec BArray< Cell_Type, Data_Type >, 31
reduced_sequence, 81	get_seq
rengine, 82	Rules < Array_Type, Data_Type >, 121
sequence, 82	
·	get_stats Model< Array Type. Data Counter Type.
set_seed, 79	-9_ 71,
simulate, 80	Data_Rule_Type, Data_Rule_Dyn_Type >,
states, 82	90
support, 82	id
update_annotations, 80	Node, 107
geese-bones.hpp	include/barry/barray-bones.hpp, 141
INITIALIZED, 171	include/barry/barray-iterator.hpp, 142
keygen_full, 172	include/barry/barray-meat-operators.hpp, 143
RULE_FUNCTION, 172	include/barry/barray-meat.hpp, 144
vec_diff, 172	include/barry/barraycell-bones.hpp, 146
vector_caster, 172	include/barry/barraycell-meat.hpp, 146
get_cell	include/barry/barraycol-bones.hpp, 147
BArray< Cell_Type, Data_Type >, 30	include/barry/barraycol-meat.hpp, 148
get_col	include/barry/barry-configuration.hpp, 148
BArray< Cell_Type, Data_Type >, 30	include/barry/barry.hpp, 150
get_col_vec	include/barry/cell-bones.hpp, 152
BArray< Cell_Type, Data_Type >, 30, 31	include/barry/cell-bottles.hpp, 152
get_counts	include/barry/col-bones.hpp, 154
Support< Array_Type, Data_Counter_Type,	include/barry/counters-bones.hpp, 154
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/counters-bones.npp, 155
133	•
get_counts_ptr	include/barry/counters/network.hpp, 155 include/barry/counters/phylo.hpp, 161
Support< Array_Type, Data_Counter_Type,	
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/model-bones.hpp, 167
133	include/barry/model-meat.hpp, 169
get_current_stats	include/barry/models/geese.hpp, 169
Support< Array_Type, Data_Counter_Type,	include/barry/models/geese/flock-bones.hpp, 170
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/models/geese/flock-meet.hpp, 170
133	include/barry/models/geese/geese-bones.hpp, 171
get_data	include/barry/models/geese/geese-meat-constructors.hpp
FreqTable $<$ T $>$, 72	173
PowerSet < Array_Type, Data_Rule_Type >, 114	include/barry/models/geese/geese-meat-likelihood.hpp,
get data ptr	173

```
include/barry/models/geese/geese-meat-likelihood exhaust.hppModel<
                                                                       Array_Type,
                                                                                       Data Counter Type,
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
include/barry/models/geese/geese-meat-predict.hpp,
                                                      likelihood
include/barry/models/geese/geese-meat-simulate.hpp,
                                                           Geese, 77
         175
                                                           Model<
                                                                                       Data Counter_Type,
                                                                       Array_Type,
include/barry/models/geese/geese-meat.hpp, 176
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
include/barry/models/geese/geese-node-bones.hpp,
                                                               90, 91
         177
                                                      likelihood
include/barry/powerset-bones.hpp, 177
                                                           model-bones.hpp, 168
include/barry/powerset-meat.hpp, 178
                                                      likelihood exhaust
include/barry/rules-bones.hpp, 179
                                                           Geese, 77
include/barry/rules-meat.hpp, 180
                                                      likelihood joint
include/barry/statscounter-bones.hpp, 181
                                                           Flock, 67
include/barry/statscounter-meat.hpp, 182
                                                      likelihood_total
include/barry/statsdb.hpp, 183
                                                           Model<
                                                                       Array_Type,
                                                                                       Data_Counter_Type,
include/barry/support-bones.hpp, 184
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
include/barry/support-meat.hpp, 185
include/barry/typedefs.hpp, 186
indices
                                                      М
    NetCounterData, 101
                                                           PowerSet < Array_Type, Data_Rule_Type >, 116
inherit_support
                                                                       Array_Type,
                                                                                       Data Counter Type,
                                                           Support<
    Geese, 77
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
init
    Counter< Array_Type, Data_Type >, 58
                                                      Мар
    Flock, 67
                                                           barry-configuration.hpp, 150
    Geese, 77
                                                      map to nodes
init_fun
                                                           Geese, 81
    Counter< Array_Type, Data_Type >, 59
                                                      MapVec_type
init node
                                                           typedefs.hpp, 188
    Geese, 77
                                                      max num elements
init_support
                                                                                       Data Counter Type,
                                                           Support<
                                                                       Array Type,
     PowerSet < Array_Type, Data_Rule_Type >, 114
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
     Support<
                 Array Type,
                                Data Counter Type,
                                                               137
         Data_Rule_Type, Data_Rule_Dyn_Type >,
                                                      Model
         133
                                                           Model<
                                                                       Array_Type,
                                                                                       Data_Counter_Type,
INITIALIZED
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
    geese-bones.hpp, 171
initialized
                                                      Model < Array_Type, Data_Counter_Type, Data_Rule_Type,
     Flock, 70
                                                               Data_Rule_Dyn_Type >, 83
    Geese, 81
                                                           \simModel, 87
insert cell
                                                           add array, 87
     BArray< Cell_Type, Data_Type >, 32
                                                           add counter, 87, 88
is_empty
                                                           add rule, 88, 89
    BArray < Cell_Type, Data_Type >, 32
                                                           add rule dyn, 89
is leaf
                                                           array frequency, 95
    Node, 106
                                                           arrays2support, 95
iter
                                                           counter_fun, 95
    ConstBArrayRowlter < Cell_Type, Data_Type >, 56
                                                           counters, 95
                                                           delete rengine, 96
keygen
                                                           first_calc_done, 96
    Model<
                Array_Type,
                                Data_Counter_Type,
                                                           get norm const, 90
         Data Rule Type, Data Rule Dyn Type >,
         96
                                                           get_pset, 90
                                                           get stats, 90
keygen default
    model-bones.hpp, 168
                                                           keygen, 96
                                                           keys2support, 96
keygen full
    geese-bones.hpp, 172
                                                           likelihood, 90, 91
                                                           likelihood total, 91
keys2support
```

Model, 86	network.hpp, 159
n_arrays_per_stats, 97	NetModel
normalizing constants, 97	network.hpp, 160
nterms, 91	NetRule
operator=, 92	network.hpp, 160
params_last, 97	NetRules
print_stats, 92	network.hpp, 160
pset_arrays, 97	NetStatsCounter
pset_probs, 98	network.hpp, 160
pset_stats, 98	NetSupport
rengine, 98	network.hpp, 160
rules, 98	Network
rules_dyn, 98	network.hpp, 160
sample, 92	Network counters, 10
•	counter_absdiff, 11
set_counters, 93 set_keygen, 93	counter_absum, 11
_ • •	
set_rengine, 93	counter_degree, 11
set_rules, 93	counter_density, 11
set_rules_dyn, 94	counter_diff, 11
set_seed, 94	counter_edges, 12
size, 94	counter_idegree, 12
size_unique, 94	counter_idegree15, 12
stats, 99	counter_isolates, 12
store_psets, 94	counter_istar2, 12
support_fun, 99	counter_mutual, 13
target_stats, 99	counter_nodecov, 13
with_pset, 99	counter_nodeicov, 13
model-bones.hpp	counter_nodematch, 13
keygen_default, 168	counter_nodeocov, 13
likelihood_, 168	counter_odegree, 14
update_normalizing_constant, 168	counter_odegree15, 14
N	counter_ostar2, 14
PowerSet < Array_Type, Data_Rule_Type >, 116	counter_ttriads, 14
Support< Array_Type, Data_Counter_Type,	NETWORK_COUNTER, 14
Data_Rule_Type, Data_Rule_Dyn_Type >,	network.hpp
138	NET_C_DATA_IDX, 158
n_arrays_per_stats	NET_C_DATA_NUM, 158
Model Array_Type, Data_Counter_Type,	NetCounter, 159
Data_Rule_Type, Data_Rule_Dyn_Type >,	NetCounters, 159
97	NetModel, 160
name	NetRule, 160 NetRules, 160
Counter< Array_Type, Data_Type >, 59	•
narray	NetStatsCounter, 160
Node, 107	NetSupport, 160
ncol	Network, 160
BArray< Cell_Type, Data_Type >, 32	NETWORK_COUNTER, 158
NET C DATA IDX	NETWORK_COUNTER_LAMBDA, 158
network.hpp, 158	NETWORK_RULE, 159
NET_C_DATA_NUM	NETWORK_RULE_LAMBDA, 159
network.hpp, 158	rules_zerodiag, 161
NetCounter	NETWORK_COUNTER
network.hpp, 159	Network counters, 14
NetCounterData, 100	network.hpp, 158
~NetCounterData, 100	NETWORK_COUNTER_LAMBDA
indices, 101	network.hpp, 158
NetCounterData, 100	NETWORK_RULE
numbers, 101	network.hpp, 159 NETWORK RULE LAMBDA
NetCounters	IVE I VVOI III_I TOLL_LAIVIDDA

network.hpp, 159	Data_Rule_Type, Data_Rule_Dyn_Type >
NetworkData, 101	91
~NetworkData, 103	ntrees
directed, 103	Flock, 69
NetworkData, 102	numbers
vertex_attr, 103	NetCounterData, 101
nfunctions	observed_counts
Flock, 70	Geese, 78
Geese, 81	offspring
nfuns	Node, 107
Flock, 68	ONE
Geese, 78	
nleafs	CHECK, 22 EXISTS, 24
Flock, 68	
Geese, 78	operator Cell_Type
nnodes	BArrayCell < Cell_Type, Data_Type >, 40
Flock, 68	BArrayCell_const< Cell_Type, Data_Type >, 43
Geese, 78	BArrayCol < Cell_Type, Data_Type >, 45
nnozero	Cell< Cell_Type >, 53
BArray< Cell_Type, Data_Type >, 33	operator!=
Node, 103	BArrayCell_const< Cell_Type, Data_Type >, 43
\sim Node, 105	BArrayCol_const< Cell_Type, Data_Type >, 48
annotations, 106	operator<
array, 106	BArrayCell_const< Cell_Type, Data_Type >, 43
arrays, 106	BArrayCol_const< Cell_Type, Data_Type >, 48
duplication, 107	operator<=
get_parent, 106	BArrayCell_const< Cell_Type, Data_Type >, 43
id, 107	BArrayCol_const< Cell_Type, Data_Type >, 48
is_leaf, 106	operator>
narray, 107	BArrayCell_const< Cell_Type, Data_Type >, 43
Node, 104, 105	BArrayCol_const< Cell_Type, Data_Type >, 49
offspring, 107	operator>=
ord, 107	BArrayCell_const< Cell_Type, Data_Type >, 44
parent, 108	BArrayCol_const< Cell_Type, Data_Type >, 49
probability, 108	operator*=
subtree_prob, 108	BArray< Cell_Type, Data_Type >, 33
visited, 108	BArrayCell< Cell_Type, Data_Type >, 40
NodeData, 109	BArrayCol < Cell_Type, Data_Type >, 46
~NodeData, 109	operator()
blengths, 110	BArray< Cell_Type, Data_Type >, 33
duplication, 110	Flock, 69
NodeData, 109	Rule < Array Type, Data Type >, 119
states, 110	Rules< Array_Type, Data_Type >, 122
nodes	vecHasher< T >, 139
	operator+=
Geese, 81 NONE	BArray< Cell_Type, Data_Type >, 33, 34
	BArrayCell< Cell_Type, Data_Type >, 40
CHECK, 22	BArrayCol< Cell_Type, Data_Type >, 46
EXISTS, 24	operator=
normalizing_constants	BArray< Cell_Type, Data_Type >, 34
Model Array_Type, Data_Counter_Type,	BArrayCell< Cell_Type, Data_Type >, 41
Data_Rule_Type, Data_Rule_Dyn_Type >,	BArrayCol Cell_Type, Data_Type >, 46
97	operator/=
nrow	BArray< Cell_Type, Data_Type >, 34
BArray< Cell_Type, Data_Type >, 33	BArrayCell< Cell_Type, Data_Type >, 34 BArrayCell< Cell_Type, Data_Type >, 41
nterms	BArrayCell Cell_Type, Data_Type >, 41 BArrayCol < Cell_Type, Data_Type >, 46
Flock, 68	
Geese, 78	operator= BArray< Cell_Type, Data_Type >, 35
Model< Array_Type, Data_Counter_Type,	BArrayCell< Cell_Type, Data_Type >, 35 BArrayCell< Cell_Type, Data_Type >, 41
	DAITAYOUI VOIL TYPE, DAIA TYPE /, 41

BArrayCol< Cell_Type, Data_Type >, 46 Cell< Cell_Type >, 53	PhyloStatsCounter, 166 PhyloSupport, 166
Counter< Array_Type, Data_Type >, 58	PHYLO_C_DATA_IDX
Counters< Array_Type, Data_Type >, 62	phylo.hpp, 163
Geese, 78, 79	PHYLO_CHECK_MISSING
Model< Array_Type, Data_Counter_Type,	phylo.hpp, 163
Data_Rule_Type, Data_Rule_Dyn_Type >,	PHYLO_COUNTER
92	phylo.hpp, 163
Rules < Array_Type, Data_Type >, 122	PHYLO_COUNTER_LAMBDA
operator==	phylo.hpp, 164
BArray< Cell_Type, Data_Type >, 35	PhyloArray
BArrayCell< Cell_Type, Data_Type >, 41	phylo.hpp, 164
BArrayCell_const< Cell_Type, Data_Type >, 43	PhyloCounter
BArrayCol < Cell_Type, Data_Type >, 47	phylo.hpp, 164
BArrayCol_const< Cell_Type, Data_Type >, 49	PhyloCounterData
operator[]	phylo.hpp, 164
Counters < Array_Type, Data_Type >, 62	PhyloCounters
PowerSet< Array_Type, Data_Rule_Type >, 115	phylo.hpp, 165
ord	PhyloModel
Node, 107	phylo.hpp, 165
out_of_range	PhyloPowerSet
BArray< Cell_Type, Data_Type >, 35	phylo.hpp, 165
	PhyloRule
params_last	phylo.hpp, 165
Model Array_Type, Data_Counter_Type,	PhyloRuleData
Data_Rule_Type, Data_Rule_Dyn_Type >,	phylo.hpp, 165
97	PhyloRules
parent	phylo.hpp, 165
Node, 108	PhyloStatsCounter
Phylo counters, 15	phylo.hpp, 166
counter_co_opt, 16	PhyloSupport
counter_cogain, 16	phylo.hpp, 166
counter_gains, 16	PowerSet
counter_gains_k_offspring, 17 counter_genes_changing, 17	PowerSet < Array_Type, Data_Rule_Type >, 112
counter_longest, 17	PowerSet < Array_Type, Data_Rule_Type >, 111
counter_loss, 17	~PowerSet, 113
counter_maxfuns, 18	add_rule, 113
counter neofun, 18	begin, 113
counter neofun a2b, 18	calc, 114
counter_overall_changes, 18	coordinates_free, 115
counter_overall_gains, 19	coordinates_locked, 115
counter_overall_loss, 19	data, 116
counter subfun, 19	EmptyArray, 116
phylo.hpp	end, 114
get_last_name, 166	get_data, 114
PHYLO_C_DATA_IDX, 163	get_data_ptr, 114
PHYLO CHECK MISSING, 163	init_support, 114
PHYLO COUNTER, 163	M, 116
PHYLO_COUNTER_LAMBDA, 164	N, 116
PhyloArray, 164	operator[], 115 PowerSet, 112
PhyloCounter, 164	
PhyloCounterData, 164	reset, 115
PhyloCounters, 165	rules, 116
PhyloModel, 165	rules_deleted, 117 size, 115
PhyloPowerSet, 165	predict
PhyloRule, 165	Geese, 79
PhyloRuleData, 165	predict_backend
PhyloRules, 165	predict_backerid

Geese, 79	Rule
print	Rule < Array_Type, Data_Type >, 118
BArray< Cell_Type, Data_Type >, 35	Rule < Array_Type, Data_Type >, 117
FreqTable $<$ T $>$, 72	\sim Rule, 118
Support< Array_Type, Data_Counter_Type,	D, 119
Data_Rule_Type, Data_Rule_Dyn_Type >,	operator(), 119
133	Rule, 118
print_observed_counts	rule_fun_default
Geese, 79	rules-bones.hpp, 180
print_stats	Rule_fun_type
Model Array_Type, Data_Counter_Type,	typedefs.hpp, 189
Data_Rule_Type, Data_Rule_Dyn_Type >, 92	RULE_FUNCTION barry.hpp, 152
probability	geese-bones.hpp, 172
Node, 108	RULE_LAMBDA
pset_arrays	barry.hpp, 152
Model< Array_Type, Data_Counter_Type,	Rules
Data_Rule_Type, Data_Rule_Dyn_Type >,	Rules< Array_Type, Data_Type >, 120
97	rules
pset_probs	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 >,	98
98	PowerSet < Array_Type, Data_Rule_Type >, 116
pset_stats	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 >,	138
98	Rules < Array_Type, Data_Type >, 119
README.md, 190	∼Rules, 120
reduced_sequence	add_rule, 121
Geese, 81	clear, 121
rengine	get_seq, 121
Flock, 70	operator(), 122
Geese, 82	operator=, 122 Rules, 120
Model< Array_Type, Data_Counter_Type,	size, 122
Data_Rule_Type, Data_Rule_Dyn_Type >,	rules-bones.hpp
98	rule_fun_default, 180
reserve	rules_deleted
BArray< Cell_Type, Data_Type >, 35	PowerSet< Array_Type, Data_Rule_Type >, 117
FreqTable $<$ T $>$, 73	rules_dyn
reset	Model< Array_Type, Data_Counter_Type,
PowerSet< Array_Type, Data_Rule_Type >, 115	Data_Rule_Type, Data_Rule_Dyn_Type >,
reset_array	98
StatsCounter < Array_Type, Data_Type >, 126	Support< Array_Type, Data_Counter_Type,
Support	$Data_Rule_Type$, $Data_Rule_Dyn_Type >$,
Data_Rule_Type, Data_Rule_Dyn_Type >,	138
134	rules_zerodiag
resize BArray < Cell_Type, Data_Type >, 36	network.hpp, 161
Entries Cell_Type >, 64	sample
rm_cell	Model Array_Type, Data_Counter_Type,
BArray< Cell_Type, Data_Type >, 36	Data_Rule_Type, Data_Rule_Dyn_Type >,
ROW	92
barray-meat-operators.hpp, 144	sequence
barray-meat.hpp, 145	Geese, 82
row	set_counters
BArray< Cell_Type, Data_Type >, 36	
Row_type	Data_Rule_Type, Data_Rule_Dyn_Type >,
typedefs.hpp, 188	93

StatsCounter< Array_Type, Data_Type >, 126	StatsCounter
Support< Array_Type, Data_Counter_Type,	StatsCounter< Array_Type, Data_Type >, 124
Data_Rule_Type, Data_Rule_Dyn_Type >,	StatsCounter< Array_Type, Data_Type >, 123
134	\sim StatsCounter, 124
set_data	add_counter, 125
BArray< Cell_Type, Data_Type >, 36	Array, 126
set_keygen	count_all, 125
Model< Array_Type, Data_Counter_Type,	count_current, 125
Data_Rule_Type, Data_Rule_Dyn_Type >,	count_init, 125
93	counter_deleted, 126
set_rengine	counters, 127
Model< Array_Type, Data_Counter_Type,	current_stats, 127
Data_Rule_Type, Data_Rule_Dyn_Type >,	EmptyArray, 127
93	reset_array, 126
set_rules	set_counters, 126
Model < Array_Type, Data_Counter_Type,	StatsCounter, 124
Data_Rule_Type, Data_Rule_Dyn_Type >,	store_psets
93	Model < Array_Type, Data_Counter_Type,
Support< Array_Type, Data_Counter_Type,	Data_Rule_Type, Data_Rule_Dyn_Type >,
Data_Rule_Type, Data_Rule_Dyn_Type >,	94
134	subtree_prob
set_rules_dyn Model < Array Type Data Counter Type	Node, 108
Model < Array_Type, Data_Counter_Type,	Support Array Type Data Counter Type
Data_Rule_Type, Data_Rule_Dyn_Type >, 94	Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,
	130
Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,	
134	support Flock, 70
	Geese, 82
set_seed Flock, 69	Support < Array_Type, Data_Counter_Type, Data_Rule_Type,
Geese, 79	Data_Rule_Dyn_Type >, 128
Model < Array_Type, Data_Counter_Type,	~Support, 131
Data_Rule_Type, Data_Rule_Dyn_Type >,	add_counter, 131
94	add_rule, 131, 132
simulate	add_rule_dyn, 132
Geese, 80	calc, 132
size	change_stats, 135
Counters < Array_Type, Data_Type >, 63	coordinates_free, 135
FreqTable $<$ T $>$, 73	coordinates_locked, 135
Model< Array_Type, Data_Counter_Type,	counters, 135
Data_Rule_Type, Data_Rule_Dyn_Type >,	current_stats, 136
94	data, 136
PowerSet < Array_Type, Data_Rule_Type >, 115	delete_counters, 136
Rules < Array_Type, Data_Type >, 122	delete_rules, 136
size_unique	delete_rules_dyn, 137
Model< Array_Type, Data_Counter_Type,	EmptyArray, 137
Data_Rule_Type, Data_Rule_Dyn_Type >,	get_counts, 133
94	get_counts_ptr, 133
source	get_current_stats, 133
Entries< Cell_Type >, 65	init_support, 133
states	M, 137
Geese, 82	max_num_elements, 137
NodeData, 110	N, 138
Statistical Models, 9	print, 133
stats Model Arrey Type Deta Counter Type	reset_array, 134
Model < Array_Type, Data_Counter_Type,	rules, 138
Data_Rule_Type, Data_Rule_Dyn_Type >, 99	rules_dyn, 138 set_counters, 134
	aci Guillela, 194

```
set_rules, 134
                                                          typedefs.hpp, 189
    set_rules_dyn, 134
                                                     vec_equal_approx
    Support, 130
                                                          typedefs.hpp, 189
support-meat.hpp
                                                     vec_inner_prod
    BARRY_SUPPORT_MEAT_HPP, 185
                                                          typedefs.hpp, 190
support_fun
                                                     vecHasher< T >, 139
    Model<
                Array Type,
                                Data Counter Type,
                                                          operator(), 139
         Data_Rule_Type, Data_Rule_Dyn_Type >,
                                                     vector caster
                                                          geese-bones.hpp, 172
swap cells
                                                     vertex_attr
    BArray< Cell_Type, Data_Type >, 37
                                                          NetworkData, 103
swap cols
                                                     visited
    BArray< Cell_Type, Data_Type >, 37
                                                          BArray< Cell_Type, Data_Type >, 39
                                                          Cell< Cell_Type >, 53
swap_rows
    BArray< Cell_Type, Data_Type >, 37
                                                          Node, 108
target
                                                     with_pset
    Entries < Cell_Type >, 65
                                                          Model<
                                                                                     Data_Counter_Type,
                                                                      Array_Type,
target_stats
                                                              Data_Rule_Type, Data_Rule_Dyn_Type >,
    Model<
                Array Type,
                                Data Counter Type,
                                                              99
         Data_Rule_Type, Data_Rule_Dyn_Type >,
                                                     zero col
                                                          BArray< Cell_Type, Data_Type >, 38
toggle_cell
                                                     zero row
    BArray< Cell_Type, Data_Type >, 37
                                                          BArray< Cell Type, Data Type >, 38
toggle lock
    BArray< Cell_Type, Data_Type >, 37
transpose
    BArray< Cell_Type, Data_Type >, 38
TWO
     CHECK, 22
    EXISTS, 24
typedefs.hpp
    Col type, 187
    Counter_fun_type, 188
    Counts_type, 188
    MapVec_type, 188
    Row type, 188
    Rule_fun_type, 189
    uint, 189
    vec_equal, 189
    vec equal approx, 189
    vec_inner_prod, 190
uint
    typedefs.hpp, 189
UKNOWN
     EXISTS, 24
update annotations
    Geese, 80
update_normalizing_constant
    model-bones.hpp, 168
val
     Entries < Cell_Type >, 65
value
    Cell< Cell_Type >, 53
vec diff
    geese-bones.hpp, 172
vec_equal
```