barry: Your go-to motif accountant 0.0-1

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1 Main Page	1
2 Module Index	3
2.1 Modules	3
3 Class Index	5
3.1 Class List	5
4 File Index	7
4.1 File List	7
5 Module Documentation	9
5.1 Counting	9
5.1.1 Detailed Description	9
5.2 Statistical Models	9
5.2.1 Detailed Description	10
5.3 Network counters	10
5.3.1 Detailed Description	10
5.3.2 Function Documentation	11
5.3.2.1 counter_absdiff()	11
5.3.2.2 counter_ctriads()	11
5.3.2.3 counter_degree()	11
5.3.2.4 counter_density()	11
5.3.2.5 counter_diff()	12
5.3.2.6 counter_edges()	12
5.3.2.7 counter_idegree()	12
5.3.2.8 counter_idegree15()	12
5.3.2.9 counter isolates()	12
5.3.2.10 counter_istar2()	13
5.3.2.11 counter_mutual()	13
5.3.2.12 counter_nodecov()	13
5.3.2.13 counter_nodeicov()	13
5.3.2.14 counter_nodematch()	13
5.3.2.15 counter_nodeocov()	14
5.3.2.16 counter_odegree()	14
5.3.2.17 counter_odegree15()	14
5.3.2.18 counter_ostar2()	14
5.3.2.19 counter_ttriads()	14
5.3.2.20 NETWORK_COUNTER()	15
5.4 Phylo counters	15
5.4.1 Detailed Description	15
5.4.2 Function Documentation	16
5.4.2 Function Documentation	16
5.4.2.2 counter_cogain()	16

	5.4.2.3 counter_gains()	1/
	5.4.2.4 counter_gains_k_offspring()	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
	5.5 Phylo rules	20
	5.5.1 Detailed Description	20
	5.5.2 Function Documentation	20
	5.5.2.1 rule_dyn_limit_changes()	20
6	Namespace Documentation	23
	6.1 barry Namespace Reference	23
	6.1.1 Detailed Description	23
	6.2 barry::counters Namespace Reference	23
	6.2.1 Detailed Description	23
	6.3 barry::counters::network Namespace Reference	24
	6.4 barry::counters::phylo Namespace Reference	24
	6.5 CHECK Namespace Reference	24
	6.5.1 Detailed Description	24
	6.5.2 Variable Documentation	24
	6.5.2.1 BOTH	24
	6.5.2.2 NONE	24
	6.5.2.3 ONE	24
	6.5.2.4 TWO	25
	6.6 EXISTS Namespace Reference	25
	6.6.1 Detailed Description	25
	6.6.2 Variable Documentation	25
	6.6.2.1 AS_ONE	25
	6.6.2.2 AS_ZERO	25
	6.6.2.3 BOTH	26
	6.6.2.4 NONE	26
	6.6.2.5 ONE	26
	6.6.2.6 TWO	26
	6.6.2.7 UKNOWN	26
7	Class Desumentation	07
1	Class Documentation	27

7.1 BArray< Cell_Type, Data_Type > Class Template Reference	27
7.1.1 Detailed Description	29
7.1.2 Constructor & Destructor Documentation	30
7.1.2.1 BArray() [1/6]	30
7.1.2.2 BArray() [2/6]	30
7.1.2.3 BArray() [3/6]	30
7.1.2.4 BArray() [4/6]	31
7.1.2.5 BArray() [5/6]	31
7.1.2.6 BArray() [6/6]	31
7.1.2.7 ~BArray()	31
7.1.3 Member Function Documentation	31
7.1.3.1 clear()	31
7.1.3.2 col()	32
7.1.3.3 D() [1/2]	32
7.1.3.4 D() [2/2]	32
7.1.3.5 default_val()	32
7.1.3.6 get_cell()	32
7.1.3.7 get_col_vec() [1/2]	32
7.1.3.8 get_col_vec() [2/2]	33
7.1.3.9 get_entries()	33
7.1.3.10 get_row_vec() [1/2]	33
7.1.3.11 get_row_vec() [2/2]	33
7.1.3.12 insert_cell() [1/3]	33
7.1.3.13 insert_cell() [2/3]	34
7.1.3.14 insert_cell() [3/3]	34
7.1.3.15 is_empty()	34
7.1.3.16 ncol()	34
7.1.3.17 nnozero()	34
7.1.3.18 nrow()	35
7.1.3.19 operator()() [1/2]	35
7.1.3.20 operator()() [2/2]	35
7.1.3.21 operator*=()	35
7.1.3.22 operator+=() [1/3]	35
7.1.3.23 operator+=() [2/3]	35
7.1.3.24 operator+=() [3/3]	36
7.1.3.25 operator-=() [1/3]	36
7.1.3.26 operator-=() [2/3]	36
7.1.3.27 operator-=() [3/3]	36
7.1.3.28 operator/=()	36
7.1.3.29 operator=() [1/2]	36
7.1.3.30 operator=() [2/2]	37
7.1.3.31 operator==()	37

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

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

7.4.3.28 operator/=()	. 56
7.4.3.29 operator=() [1/2]	. 56
7.4.3.30 operator=() [2/2]	. 57
7.4.3.31 operator==()	. 57
7.4.3.32 out_of_range()	. 57
7.4.3.33 print()	. 57
7.4.3.34 reserve()	. 57
7.4.3.35 resize()	. 58
7.4.3.36 rm_cell()	. 58
7.4.3.37 row()	. 58
7.4.3.38 set_data()	. 58
7.4.3.39 swap_cells()	. 59
7.4.3.40 swap_cols()	. 59
7.4.3.41 swap_rows()	. 59
7.4.3.42 toggle_cell()	. 59
7.4.3.43 toggle_lock()	. 60
7.4.3.44 transpose()	. 60
7.4.3.45 zero_col()	. 60
7.4.3.46 zero_row()	. 60
7.4.4 Friends And Related Function Documentation	. 60
7.4.4.1 BArrayCell< Cell_Type, Data_Type >	. 60
7.4.4.2 BArrayCell_const< Cell_Type, Data_Type >	. 61
7.4.5 Member Data Documentation	. 61
7.4.5.1 visited	. 61
7.5 BArrayDenseCell< Cell_Type, Data_Type > Class Template Reference	. 61
7.5.1 Detailed Description	. 62
7.5.2 Constructor & Destructor Documentation	. 62
7.5.2.1 BArrayDenseCell()	. 62
7.5.2.2 ~BArrayDenseCell()	. 62
7.5.3 Member Function Documentation	. 62
7.5.3.1 operator Cell_Type()	. 62
7.5.3.2 operator*=()	. 63
7.5.3.3 operator+=()	. 63
7.5.3.4 operator-=()	. 63
7.5.3.5 operator/=()	. 63
7.5.3.6 operator=()	. 63
7.5.3.7 operator==()	. 64
7.6 BArrayDenseCell_const< Cell_Type, Data_Type > Class Template Reference	. 64
7.6.1 Detailed Description	. 64
7.6.2 Constructor & Destructor Documentation	. 64
7.6.2.1 BArrayDenseCell_const()	. 65
7.6.2.2 ~BArrayDenseCell_const()	. 65

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

7.8.3.11 operator>=()	74
7.8.3.12 size()	74
7.9 Cell< Cell_Type > Class Template Reference	74
7.9.1 Detailed Description	75
7.9.2 Constructor & Destructor Documentation	75
7.9.2.1 Cell() [1/7]	75
7.9.2.2 Cell() [2/7]	75
7.9.2.3 ~Cell()	75
7.9.2.4 Cell() [3/7]	76
7.9.2.5 Cell() [4/7]	76
7.9.2.6 Cell() [5/7]	76
7.9.2.7 Cell() [6/7]	76
7.9.2.8 Cell() [7/7]	76
7.9.3 Member Function Documentation	76
7.9.3.1 add() [1/4]	77
7.9.3.2 add() [2/4]	77
7.9.3.3 add() [3/4]	77
7.9.3.4 add() [4/4]	77
7.9.3.5 operator Cell_Type()	77
7.9.3.6 operator"!=()	77
7.9.3.7 operator=() [1/2]	78
7.9.3.8 operator=() [2/2]	78
7.9.3.9 operator==()	78
7.9.4 Member Data Documentation	78
7.9.4.1 value	78
7.9.4.2 visited	78
$7.10 \ ConstBArray Rowlter < Cell_Type, \ Data_Type > Class \ Template \ Reference \\ \ \dots \dots$	79
7.10.1 Detailed Description	79
7.10.2 Constructor & Destructor Documentation	79
7.10.2.1 ConstBArrayRowlter()	80
7.10.2.2 ~ConstBArrayRowlter()	80
7.10.3 Member Data Documentation	80
7.10.3.1 Array	80
7.10.3.2 current_col	80
7.10.3.3 current_row	80
7.10.3.4 iter	81
7.11 Counter< Array_Type, Data_Type > Class Template Reference	81
7.11.1 Detailed Description	82
7.11.2 Constructor & Destructor Documentation	82
7.11.2.1 Counter() [1/4]	82
7.11.2.2 Counter() [2/4]	82
7.11.2.3 Counter() [3/4]	83

7.11.2.4 Counter() [4/4]	83
7.11.2.5 ~Counter()	83
7.11.3 Member Function Documentation	83
7.11.3.1 count()	83
7.11.3.2 init()	84
7.11.3.3 operator=() [1/2]	84
7.11.3.4 operator=() [2/2]	84
7.11.4 Member Data Documentation	84
7.11.4.1 count_fun	84
7.11.4.2 data	85
7.11.4.3 delete_data	85
7.11.4.4 desc	85
7.11.4.5 init_fun	85
7.11.4.6 name	85
7.12 Counters < Array_Type, Data_Type > Class Template Reference	86
7.12.1 Detailed Description	86
7.12.2 Constructor & Destructor Documentation	86
7.12.2.1 Counters() [1/3]	87
7.12.2.2 ~Counters()	87
7.12.2.3 Counters() [2/3]	87
7.12.2.4 Counters() [3/3]	87
7.12.3 Member Function Documentation	88
7.12.3.1 add_counter() [1/3]	88
7.12.3.2 add_counter() [2/3]	88
7.12.3.3 add_counter() [3/3]	88
7.12.3.4 clear()	88
7.12.3.5 operator=() [1/2]	88
7.12.3.6 operator=() [2/2]	89
7.12.3.7 operator[]()	89
7.12.3.8 size()	90
7.13 Entries < Cell_Type > Class Template Reference	90
7.13.1 Detailed Description	90
7.13.2 Constructor & Destructor Documentation	91
7.13.2.1 Entries() [1/2]	91
7.13.2.2 Entries() [2/2]	91
7.13.2.3 ~Entries()	91
7.13.3 Member Function Documentation	91
7.13.3.1 resize()	91
7.13.4 Member Data Documentation	92
7.13.4.1 source	92
7.13.4.2 target	92
7.13.4.3 val	92

7.14 Flock Class Reference	92
7.14.1 Detailed Description	93
7.14.2 Constructor & Destructor Documentation	93
7.14.2.1 Flock()	94
7.14.2.2 ~Flock()	94
7.14.3 Member Function Documentation	94
7.14.3.1 add_data()	94
7.14.3.2 get_counters()	94
7.14.3.3 get_support()	95
7.14.3.4 init()	95
7.14.3.5 likelihood_joint()	95
7.14.3.6 nfuns()	95
7.14.3.7 nleafs()	96
7.14.3.8 nnodes()	96
7.14.3.9 nterms()	96
7.14.3.10 ntrees()	96
7.14.3.11 operator()()	96
7.14.3.12 set_seed()	97
7.14.3.13 support_size()	97
7.14.4 Member Data Documentation	97
7.14.4.1 dat	97
7.14.4.2 initialized	97
7.14.4.3 model	98
7.14.4.4 nfunctions	98
7.14.4.5 rengine	98
7.15 FreqTable $<$ T $>$ Class Template Reference	98
7.15.1 Detailed Description	99
7.15.2 Constructor & Destructor Documentation	99
7.15.2.1 FreqTable()	99
7.15.2.2 ~FreqTable()	99
7.15.3 Member Function Documentation	99
7.15.3.1 add()	99
7.15.3.2 as_vector()	100
7.15.3.3 clear()	100
7.15.3.4 get_data()	100
7.15.3.5 get_data_ptr()	100
7.15.3.6 print()	100
7.15.3.7 reserve()	101
7.15.3.8 size()	101
7.16 Geese Class Reference	101
7.16.1 Detailed Description	103
7.16.2 Constructor & Destructor Documentation	103

7.16.2.1 Geese() [1/4]
7.16.2.2 Geese() [2/4]
7.16.2.3 Geese() [3/4]
7.16.2.4 Geese() [4/4]
7.16.2.5 ~Geese()
7.16.3 Member Function Documentation
7.16.3.1 calc_reduced_sequence()
7.16.3.2 calc_sequence()
7.16.3.3 get_counters()
7.16.3.4 get_model()
7.16.3.5 get_probabilities()
7.16.3.6 get_rengine()
7.16.3.7 get_states()
7.16.3.8 get_support()
7.16.3.9 inherit_support()
7.16.3.10 init()
7.16.3.11 init_node()
7.16.3.12 likelihood()
7.16.3.13 likelihood_exhaust()
7.16.3.14 nfuns()
7.16.3.15 nleafs()
7.16.3.16 nnodes()
7.16.3.17 nterms()
7.16.3.18 observed_counts()
7.16.3.19 operator=() [1/2]
7.16.3.20 operator=() [2/2]
7.16.3.21 parse_polytomies()
7.16.3.22 predict()
7.16.3.23 predict_backend()
7.16.3.24 print_observed_counts()
7.16.3.25 set_seed()
7.16.3.26 simulate()
7.16.3.27 support_size()
7.16.3.28 update_annotations()
7.16.4 Member Data Documentation
7.16.4.1 delete_rengine
7.16.4.2 delete_support
7.16.4.3 initialized
7.16.4.4 map_to_nodes
7.16.4.5 nfunctions
7.16.4.6 nodes
7.16.4.7 reduced sequence 11

	7.16.4.8 sequence
	I< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class Tem-Reference
7.17	1 Detailed Description
7.17	2 Constructor & Destructor Documentation
	7.17.2.1 Model() [1/3]
	7.17.2.2 Model() [2/3]
	7.17.2.3 Model() [3/3]
	7.17.2.4 ~Model()
7.17	3 Member Function Documentation
	7.17.3.1 add_array()
	7.17.3.2 add_counter() [1/3]
	7.17.3.3 add_counter() [2/3]
	7.17.3.4 add_counter() [3/3]
	7.17.3.5 add_rule() [1/3]
	7.17.3.6 add_rule() [2/3]
	7.17.3.7 add_rule() [3/3]
	7.17.3.8 add_rule_dyn() [1/3]
	7.17.3.9 add_rule_dyn() [2/3]
	7.17.3.10 add_rule_dyn() [3/3]
	7.17.3.11 get_counters()
	7.17.3.12 get_norm_const()
	7.17.3.13 get_pset()
	7.17.3.14 get_rengine()
	7.17.3.15 get_rules()
	7.17.3.16 get_rules_dyn()
	7.17.3.17 get_stats()
	7.17.3.18 get_support()
	7.17.3.19 likelihood() [1/3]
	7.17.3.20 likelihood() [2/3]
	7.17.3.21 likelihood() [3/3]
	7.17.3.22 likelihood_total()
	7.17.3.23 nterms()
	7.17.3.24 operator=()
	7.17.3.25 print_stats()
	7.17.3.26 sample() [1/2]
	7.17.3.27 sample() [2/2]
	7.17.3.28 set_counters()
	7.17.3.29 set_keygen()
	7.17.3.30 set_rengine()
	7.17.3.31 set_rules()
	7.17.3.32 set_rules_dyn()

7.17.3.33 set_seed()	2
7.17.3.34 size()	2
7.17.3.35 size_unique()	23
7.17.3.36 store_psets()	23
7.17.3.37 support_size()	:3
7.18 NetCounterData Class Reference	:3
7.18.1 Detailed Description	4
7.18.2 Constructor & Destructor Documentation	:4
7.18.2.1 NetCounterData() [1/2]	:4
7.18.2.2 NetCounterData() [2/2]	:4
7.18.2.3 ~NetCounterData()	4
7.18.3 Member Data Documentation	:4
7.18.3.1 indices	:5
7.18.3.2 numbers	:5
7.19 NetworkData Class Reference	:5
7.19.1 Detailed Description	:5
7.19.2 Constructor & Destructor Documentation	:6
7.19.2.1 NetworkData() [1/3]	6
7.19.2.2 NetworkData() [2/3]	6
7.19.2.3 NetworkData() [3/3]	:6
7.19.2.4 ~NetworkData()	:7
7.19.3 Member Data Documentation	:7
7.19.3.1 directed	:7
7.19.3.2 vertex_attr	:7
7.20 Node Class Reference	:7
7.20.1 Detailed Description	8
7.20.2 Constructor & Destructor Documentation	8:
7.20.2.1 Node() [1/5]	:9
7.20.2.2 Node() [2/5]	:9
7.20.2.3 Node() [3/5]	9
7.20.2.4 Node() [4/5]	:9
7.20.2.5 Node() [5/5]	:9
7.20.2.6 ∼Node()	0
7.20.3 Member Function Documentation	0
7.20.3.1 get_parent()	0
7.20.3.2 is_leaf()	0
7.20.3.3 noffspring()	0
7.20.4 Member Data Documentation	0
7.20.4.1 annotations	0
7.20.4.2 array	1
7.20.4.3 arrays	1
7.20.4.4 duplication	1

7.20.4.5 id	131
7.20.4.6 narray	131
7.20.4.7 offspring	132
7.20.4.8 ord	132
7.20.4.9 parent	132
7.20.4.10 probability	132
7.20.4.11 subtree_prob	132
7.20.4.12 visited	133
7.21 NodeData Class Reference	133
7.21.1 Detailed Description	133
7.21.2 Constructor & Destructor Documentation	133
7.21.2.1 NodeData() [1/2]	134
7.21.2.2 NodeData() [2/2]	134
7.21.2.3 ~NodeData()	134
7.21.3 Member Data Documentation	134
7.21.3.1 blengths	134
7.21.3.2 duplication	134
7.21.3.3 states	135
7.22 PhyloRuleDynData Class Reference	135
7.22.1 Detailed Description	135
7.22.2 Constructor & Destructor Documentation	135
7.22.2.1 PhyloRuleDynData()	135
$7.22.2.2 \sim$ PhyloRuleDynData()	136
7.22.3 Member Data Documentation	136
7.22.3.1 counts	136
7.22.3.2 duplication	136
7.22.3.3 lb	136
7.22.3.4 pos	136
7.22.3.5 ub	136
7.23 PowerSet< Array_Type, Data_Rule_Type > Class Template Reference	137
7.23.1 Detailed Description	138
7.23.2 Constructor & Destructor Documentation	138
7.23.2.1 PowerSet() [1/3]	138
7.23.2.2 PowerSet() [2/3]	138
7.23.2.3 PowerSet() [3/3]	139
7.23.2.4 ~PowerSet()	139
7.23.3 Member Function Documentation	139
7.23.3.1 add_rule() [1/3]	139
7.23.3.2 add_rule() [2/3]	139
7.23.3.3 add_rule() [3/3]	139
7.23.3.4 begin()	140
7.23.3.5 calc()	140

7.23.3.6 end()	0
7.23.3.7 get_data()	-0
7.23.3.8 get_data_ptr()	0
7.23.3.9 init_support()	1
7.23.3.10 operator[]()	1
7.23.3.11 reset()	1
7.23.3.12 size()	1
7.23.4 Member Data Documentation	1
7.23.4.1 coordinates_free	1
7.23.4.2 coordinates_locked	-2
7.23.4.3 data	2
7.23.4.4 EmptyArray	2
7.23.4.5 M	2
7.23.4.6 N	2
7.23.4.7 rules	3
7.23.4.8 rules_deleted	3
7.24 Rule < Array_Type, Data_Type > Class Template Reference	3
7.24.1 Detailed Description	4
7.24.2 Constructor & Destructor Documentation	4
7.24.2.1 Rule() [1/2]	4
7.24.2.2 Rule() [2/2]	4
7.24.2.3 ~Rule()	4
7.24.3 Member Function Documentation	-5
7.24.3.1 D()	-5
7.24.3.2 operator()()	-5
7.25 Rules < Array_Type, Data_Type > Class Template Reference	-5
7.25.1 Detailed Description	-6
7.25.2 Constructor & Destructor Documentation	-6
7.25.2.1 Rules() [1/2]	-6
7.25.2.2 Rules() [2/2]	-6
7.25.2.3 ~Rules()	7
7.25.3 Member Function Documentation	7
7.25.3.1 add_rule() [1/3]	7
7.25.3.2 add_rule() [2/3]	7
7.25.3.3 add_rule() [3/3]	7
7.25.3.4 clear()	7
7.25.3.5 get_seq()	7
7.25.3.6 operator()()	8
7.25.3.7 operator=()	8
7.25.3.8 size()	9
7.26 StatsCounter< Array_Type, Data_Type > Class Template Reference	9
7.26.1 Detailed Description	.9

7.26.2 Constructor & Destructor Documentation	150
7.26.2.1 StatsCounter() [1/2]	150
7.26.2.2 StatsCounter() [2/2]	150
7.26.2.3 ~StatsCounter()	150
7.26.3 Member Function Documentation	150
7.26.3.1 add_counter() [1/2]	150
7.26.3.2 add_counter() [2/2]	. 151
7.26.3.3 count_all()	151
7.26.3.4 count_current()	151
7.26.3.5 count_init()	151
7.26.3.6 get_counters()	151
7.26.3.7 reset_array()	151
7.26.3.8 set_counters()	152
$7.27 \; Support < Array_Type, \; Data_Counter_Type, \; Data_Rule_Type, \; Data_Rule_Dyn_Type > Class \; Temporal Counter_Type, \; Data_Rule_Dyn_Type > Class \; Temporal Counter_Type, \; Data_Rule_Dyn_Type > Class \; Temporal Counter_Type, \; Data_Rule_Dyn_Type > Class \; Temporal Counter_Dyn_Type > Class \; Temporal Counter_Dyn_Dyn_Dyn_Dyn_Dyn_Dyn_Dyn_Dyn_Dyn_Dyn$	
plate Reference	
7.27.1 Detailed Description	
7.27.2 Constructor & Destructor Documentation	
7.27.2.1 Support() [1/3]	
7.27.2.2 Support() [2/3]	
7.27.2.3 Support() [3/3]	
7.27.2.4 ~Support()	
7.27.3 Member Function Documentation	
7.27.3.1 add_counter() [1/2]	
7.27.3.2 add_counter() [2/2]	
7.27.3.3 add_rule() [1/2]	
7.27.3.4 add_rule() [2/2]	
7.27.3.5 add_rule_dyn() [1/2]	
7.27.3.6 add_rule_dyn() [2/2]	. 157
7.27.3.7 calc()	
7.27.3.8 get_counters()	158
7.27.3.9 get_counts()	158
7.27.3.10 get_counts_ptr()	
7.27.3.11 get_current_stats()	158
7.27.3.12 get_data()	159
7.27.3.13 get_rules()	159
7.27.3.14 get_rules_dyn()	159
7.27.3.15 init_support()	159
7.27.3.16 print()	160
7.27.3.17 reset_array() [1/2]	
7.27.3.18 reset_array() [2/2]	160
7.27.3.19 set_counters()	160
7.27.3.20 set_rules()	160

	7.27.3.21 set_rules_dyn()	161
	7.27.4 Member Data Documentation	161
	7.27.4.1 change_stats	161
	7.27.4.2 coordinates_free	161
	7.27.4.3 coordinates_locked	161
	7.27.4.4 current_stats	162
	7.27.4.5 delete_counters	162
	7.27.4.6 delete_rules	162
	7.27.4.7 delete_rules_dyn	162
	7.27.4.8 M	162
	7.27.4.9 max_num_elements	163
	7.27.4.10 N	163
	7.28 vecHasher $<$ T $>$ Struct Template Reference	163
	7.28.1 Detailed Description	163
	7.28.2 Member Function Documentation	163
	7.28.2.1 operator()()	163
۰.	Tile Desumentation	105
8 1	File Documentation	165
	8.1 include/barry/barray-bones.hpp File Reference	
	8.1.1 Macro Definition Documentation	
	8.1.1.1 BARRAY_BONES_HPP	
	8.2 include/barry/barray-iterator.hpp File Reference	
	8.3 include/barry/barray-meat-operators.hpp File Reference	
	8.3.1 Macro Definition Documentation	
	8.3.1.1 BARRY_BARRAY_MEAT_OPERATORS_HPP	
	8.3.1.2 COL	
	8.3.1.3 ROW	
	8.3.2 Function Documentation	
	8.3.2.1 checkdim_()	
	8.4 include/barry/barray-meat.hpp File Reference	
	8.4.1 Macro Definition Documentation	
	8.4.1.1 COL	
	8.4.1.2 ROW	
	8.5 include/barry/barraycell-bones.hpp File Reference	
	8.6 include/barry/barraycell-meat.hpp File Reference	
	8.7 include/barry/barraydense-bones.hpp File Reference	
	8.8 include/barry/barraydense-meet.hpp File Reference	
	8.8.1 Macro Definition Documentation	
	8.8.1.1 BARRY_BARRAYDENSE_MEAT_HPP	
	8.8.1.2 COL	
	8.8.1.3 POS	
	8.8.1.4 ROW	175

8.8.1.5 ZERO_CELL
8.9 include/barry/barraydensecell-bones.hpp File Reference
8.10 include/barry/barraydensecell-meat.hpp File Reference
8.10.1 Macro Definition Documentation
8.10.1.1 BARRY_BARRAYDENSECELL_MEAT_HPP
8.10.1.2 POS
8.11 include/barry/barrayvector-bones.hpp File Reference
8.12 include/barry/barrayvector-meat.hpp File Reference
8.12.1 Macro Definition Documentation
8.12.1.1 BARRY_BARRAYVECTOR_MEAT_HPP
8.13 include/barry/barry-configuration.hpp File Reference
8.13.1 Macro Definition Documentation
8.13.1.1 BARRY_CHECK_SUPPORT
8.13.1.2 BARRY_ISFINITE
8.13.1.3 BARRY_MAX_NUM_ELEMENTS
8.13.1.4 BARRY_SAFE_EXP
8.13.1.5 printf_barry
8.13.2 Typedef Documentation
8.13.2.1 Map
8.14 include/barry/barry.hpp File Reference
8.14.1 Macro Definition Documentation
8.14.1.1 BARRY_HPP
8.14.1.2 BARRY_VERSION
8.14.1.3 COUNTER_FUNCTION
8.14.1.4 COUNTER_LAMBDA
8.14.1.5 RULE_FUNCTION
8.14.1.6 RULE_LAMBDA
8.15 include/barry/cell-bones.hpp File Reference
8.16 include/barry/cell-meat.hpp File Reference
8.17 include/barry/col-bones.hpp File Reference
8.18 include/barry/counters-bones.hpp File Reference
8.19 include/barry/counters-meat.hpp File Reference
8.20 include/barry/counters/network.hpp File Reference
8.20.1 Macro Definition Documentation
8.20.1.1 NET_C_DATA_IDX
8.20.1.2 NET_C_DATA_NUM
8.20.1.3 NETWORK_COUNTER
8.20.1.4 NETWORK_COUNTER_LAMBDA
8.20.1.5 NETWORK_RULE
8.20.1.6 NETWORK_RULE_LAMBDA
8.20.2 Typedef Documentation
8 20 2 1 NetCounter

8.20.2.2 NetCounters
8.20.2.3 NetModel
8.20.2.4 NetRule
8.20.2.5 NetRules
8.20.2.6 NetStatsCounter
8.20.2.7 NetSupport
8.20.2.8 Network
8.20.3 Function Documentation
8.20.3.1 rules_zerodiag()
8.21 include/barry/counters/phylo.hpp File Reference
8.21.1 Macro Definition Documentation
8.21.1.1 PHYLO_CHECK_MISSING
8.21.1.2 PHYLO_COUNTER_LAMBDA
8.21.1.3 PHYLO_RULE_DYN_LAMBDA
8.21.2 Typedef Documentation
8.21.2.1 PhyloArray
8.21.2.2 PhyloCounter
8.21.2.3 PhyloCounterData
8.21.2.4 PhyloCounters
8.21.2.5 PhyloModel
8.21.2.6 PhyloPowerSet
8.21.2.7 PhyloRule
8.21.2.8 PhyloRuleData
8.21.2.9 PhyloRuleDyn
8.21.2.10 PhyloRules
8.21.2.11 PhyloRulesDyn
8.21.2.12 PhyloStatsCounter
8.21.2.13 PhyloSupport
8.21.3 Function Documentation
8.21.3.1 get_last_name()
8.22 include/barry/model-bones.hpp File Reference
8.22.1 Function Documentation
8.22.1.1 keygen_default()
8.22.1.2 likelihood_()
8.22.1.3 update_normalizing_constant()
8.23 include/barry/model-meat.hpp File Reference
8.24 include/barry/models/geese.hpp File Reference
8.25 include/barry/models/geese/flock-bones.hpp File Reference
8.26 include/barry/models/geese/flock-meet.hpp File Reference
8.27 include/barry/models/geese/geese-bones.hpp File Reference
8.27.1 Macro Definition Documentation
8 27 1 1 INITIAL IZED 20.

Index

8.27.2 Function Documentation	204
8.27.2.1 keygen_full()	204
8.27.2.2 RULE_FUNCTION()	204
8.27.2.3 vec_diff()	204
8.27.2.4 vector_caster()	204
8.28 include/barry/models/geese/geese-meat-constructors.hpp File Reference	205
8.29 include/barry/models/geese/geese-meat-likelihood.hpp File Reference	205
8.30 include/barry/models/geese/geese-meat-likelihood_exhaust.hpp File Reference	206
8.31 include/barry/models/geese/geese-meat-predict.hpp File Reference	207
8.32 include/barry/models/geese/geese-meat-simulate.hpp File Reference	207
8.33 include/barry/models/geese/geese-meat.hpp File Reference	208
8.34 include/barry/models/geese/geese-node-bones.hpp File Reference	208
8.35 include/barry/powerset-bones.hpp File Reference	209
8.36 include/barry/powerset-meat.hpp File Reference	210
8.37 include/barry/rules-bones.hpp File Reference	212
8.37.1 Function Documentation	213
8.37.1.1 rule_fun_default()	213
8.38 include/barry/rules-meat.hpp File Reference	213
8.39 include/barry/statscounter-bones.hpp File Reference	214
8.40 include/barry/statscounter-meat.hpp File Reference	215
8.41 include/barry/statsdb.hpp File Reference	216
8.42 include/barry/support-bones.hpp File Reference	217
8.43 include/barry/support-meat.hpp File Reference	219
8.43.1 Macro Definition Documentation	220
8.43.1.1 BARRY_SUPPORT_MEAT_HPP	220
8.44 include/barry/typedefs.hpp File Reference	220
8.44.1 Typedef Documentation	222
8.44.1.1 Col_type	222
8.44.1.2 Counter_fun_type	222
8.44.1.3 Counts_type	222
8.44.1.4 MapVec_type	222
8.44.1.5 Row_type	223
8.44.1.6 Rule_fun_type	223
8.44.1.7 uint	223
8.44.2 Function Documentation	223
8.44.2.1 vec_equal()	223
8.44.2.2 vec_equal_approx()	224
8.44.2.3 vec_inner_prod()	224
8.45 README.md File Reference	224

225

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

4 Module Index

Class Index

3.1 Class List

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

BArray< Cell_Type, Data_Type >	
Baseline class for binary arrays	27
BArrayCell< Cell_Type, Data_Type >	41
BArrayCell_const< Cell_Type, Data_Type >	43
BArrayDense < Cell_Type, Data_Type >	
Baseline class for binary arrays	46
BArrayDenseCell< Cell_Type, Data_Type >	61
BArrayDenseCell_const< Cell_Type, Data_Type >	64
BArrayVector< Cell_Type, Data_Type >	
Row or column of a BArray	66
BArrayVector_const< Cell_Type, Data_Type >	70
Cell< Cell_Type >	
Entries in BArray. For now, it only has two members:	74
ConstBArrayRowlter< Cell_Type, Data_Type >	79
Counter< Array_Type, Data_Type >	
A counter function based on change statistics	81
Counters < Array_Type, Data_Type >	
Vector of counters	86
Entries < Cell_Type >	
A wrapper class to store source, target, val from a BArray object	90
Flock	
A Flock is a group of Geese	92
FreqTable < T >	
Database of statistics	98
Geese	
Annotated Phylo Model	101
Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >	
General framework for discrete exponential models. This class allows generating discrete expo-	
·	111
NetCounterData	
Data class used to store arbitrary uint or double vectors	123
NetworkData	
Data class for Networks	125
Node	
A single node for the model	127

6 Class Index

NodeData	
Data definition for the PhyloArray class	133
PhyloRuleDynData	
PowerSet< Array_Type, Data_Rule_Type >	
Powerset of a binary array	37
Rule < Array_Type, Data_Type >	
Rule for determining if a cell should be included in a sequence	43
Rules < Array_Type, Data_Type >	
Vector of objects of class Rule	45
StatsCounter< Array_Type, Data_Type >	
Count stats for a single Array	49
Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >	
Compute the support of sufficient statistics	152
vecHasher< T >	163

File Index

4.1 File List

Here is a list of all files with brief descriptions:

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

8 File Index

include/barry/models/geese/flock-meet.hpp	02
include/barry/models/geese/geese-bones.hpp	03
include/barry/models/geese/geese-meat-constructors.hpp	ე5
include/barry/models/geese/geese-meat-likelihood.hpp	ე5
include/barry/models/geese/geese-meat-likelihood_exhaust.hpp	ე6
include/barry/models/geese/geese-meat-predict.hpp	ე7
include/barry/models/geese/geese-meat-simulate.hpp	ე7
include/barry/models/geese/geese-meat.hpp	3 80
include/barry/models/geese/geese-node-bones.hpp	ე8

Module Documentation

5.1 Counting

Classes

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

Data definition for the PhyloArray class.

class Counter< Array_Type, Data_Type >

A counter function based on change statistics.

5.1.1 Detailed Description

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

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

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

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

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

5.2 Statistical Models

Statistical models available in barry.

10 Module Documentation

Classes

class Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >

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

· class Flock

A Flock is a group of Geese.

· class Geese

Annotated Phylo Model.

5.2.1 Detailed Description

Statistical models available in barry.

5.3 Network counters

Counters for network models.

Functions

• void counter_edges (NetCounters *counters)

Number of edges.

void counter isolates (NetCounters *counters)

Number of isolated vertices.

void counter_mutual (NetCounters *counters)

Number of mutual ties.

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

Sum of absolute attribute difference between ego and alter.

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

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

- NETWORK_COUNTER (init_single_attr)
- void counter nodeicov (NetCounters *counters, uint attr id)
- void counter_nodeocov (NetCounters *counters, uint attr_id)
- void counter_nodecov (NetCounters *counters, uint attr_id)
- void counter_nodematch (NetCounters *counters, uint attr_id)
- void counter_idegree (NetCounters *counters, std::vector< uint > d)

Counts number of vertices with a given in-degree.

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

Counts number of vertices with a given out-degree.

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

Counts number of vertices with a given out-degree.

5.3.1 Detailed Description

Counters for network models.

5.3 Network counters

Parameters

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

5.3.2 Function Documentation

5.3.2.1 counter_absdiff()

Sum of absolute attribute difference between ego and alter.

Definition at line 419 of file network.hpp.

5.3.2.2 counter_ctriads()

Definition at line 322 of file network.hpp.

5.3.2.3 counter_degree()

Counts number of vertices with a given out-degree.

Definition at line 690 of file network.hpp.

5.3.2.4 counter_density()

Definition at line 361 of file network.hpp.

12 Module Documentation

5.3.2.5 counter_diff()

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

Definition at line 461 of file network.hpp.

5.3.2.6 counter_edges()

Number of edges.

Definition at line 128 of file network.hpp.

5.3.2.7 counter_idegree()

Counts number of vertices with a given in-degree.

Definition at line 604 of file network.hpp.

5.3.2.8 counter_idegree15()

Definition at line 377 of file network.hpp.

5.3.2.9 counter_isolates()

Number of isolated vertices.

Definition at line 142 of file network.hpp.

5.3 Network counters

5.3.2.10 counter_istar2()

Definition at line 210 of file network.hpp.

5.3.2.11 counter_mutual()

Number of mutual ties.

Definition at line 172 of file network.hpp.

5.3.2.12 counter_nodecov()

Definition at line 558 of file network.hpp.

5.3.2.13 counter_nodeicov()

Definition at line 520 of file network.hpp.

5.3.2.14 counter_nodematch()

Definition at line 578 of file network.hpp.

14 Module Documentation

5.3.2.15 counter_nodeocov()

Definition at line 539 of file network.hpp.

5.3.2.16 counter_odegree()

Counts number of vertices with a given out-degree.

Definition at line 646 of file network.hpp.

5.3.2.17 counter_odegree15()

Definition at line 397 of file network.hpp.

5.3.2.18 counter_ostar2()

Definition at line 228 of file network.hpp.

5.3.2.19 counter_ttriads()

Definition at line 247 of file network.hpp.

5.4 Phylo counters 15

5.3.2.20 NETWORK_COUNTER()

Definition at line 503 of file network.hpp.

5.4 Phylo counters

Counters for phylogenetic modeling.

Functions

- void counter_overall_gains (PhyloCounters *counters, bool duplication=true)
 Overall functional gains.
- void counter_gains (PhyloCounters *counters, std::vector< uint > nfun, bool duplication=true)
 Functional gains for a specific function (nfun).
- void counter_gains_k_offspring (PhyloCounters *counters, std::vector< uint > nfun, uint k=1u, bool duplication=true)

k genes gain function nfun

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

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

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

Overall functional loss.

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

Total count of losses for an specific function.

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

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

- void counter_subfun (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)
 - Total count of Sub-functionalization events.
- void counter_cogain (PhyloCounters *counters, uint nfunA, uint nfunB, bool duplication=true)

Co-evolution (joint gain or loss)

void counter_longest (PhyloCounters *counters)

Longest branch mutates (either by gain or by loss)

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

Total number of neofunctionalization events.

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

Total number of neofunctionalization events.

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

5.4.1 Detailed Description

Counters for phylogenetic modeling.

16 Module Documentation

Parameters

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

5.4.2 Function Documentation

5.4.2.1 counter_co_opt()

Function co-opting.

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

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

This algorithm implements the change statistic.

Definition at line 1084 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 714 of file phylo.hpp.

5.4 Phylo counters

5.4.2.3 counter_gains()

Functional gains for a specific function (nfun).

Definition at line 152 of file phylo.hpp.

5.4.2.4 counter_gains_k_offspring()

k genes gain function nfun

Definition at line 194 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 268 of file phylo.hpp.

5.4.2.6 counter_longest()

Longest branch mutates (either by gain or by loss)

Definition at line 773 of file phylo.hpp.

18 Module Documentation

5.4.2.7 counter_loss()

Total count of losses for an specific function.

Definition at line 515 of file phylo.hpp.

5.4.2.8 counter_maxfuns()

Cap the number of functions per gene.

Definition at line 431 of file phylo.hpp.

5.4.2.9 counter_neofun()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 878 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 963 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 562 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 112 of file phylo.hpp.

5.4.2.13 counter_overall_loss()

Overall functional loss.

Definition at line 385 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 628 of file phylo.hpp.

20 Module Documentation

5.5 Phylo rules

Rules for phylogenetic modeling.

Classes

• class PhyloRuleDynData

Functions

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

Overall functional gains.

5.5.1 Detailed Description

Rules for phylogenetic modeling.

Parameters

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

5.5.2 Function Documentation

5.5.2.1 rule_dyn_limit_changes()

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

Overall functional gains.

Parameters

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

5.5 Phylo rules 21

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

Definition at line 1217 of file phylo.hpp.

22 Module Documentation

Chapter 6

Namespace Documentation

6.1 barry Namespace Reference

barry: Your go-to motif accountant

Namespaces

counters

Tree class and Treelterator class.

6.1.1 Detailed Description

barry: Your go-to motif accountant

6.2 barry::counters Namespace Reference

Tree class and Treelterator class.

Namespaces

- network
- phylo

6.2.1 Detailed Description

Tree class and Treelterator class.

6.3 barry::counters::network Namespace Reference

6.4 barry::counters::phylo Namespace Reference

6.5 CHECK Namespace Reference

Integer constants used to specify which cell should be check.

Variables

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

6.5.1 Detailed Description

Integer constants used to specify which cell should be check.

6.5.2 Variable Documentation

6.5.2.1 BOTH

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

Definition at line 20 of file typedefs.hpp.

6.5.2.2 NONE

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

Definition at line 21 of file typedefs.hpp.

6.5.2.3 ONE

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

Definition at line 22 of file typedefs.hpp.

6.5.2.4 TWO

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

Definition at line 23 of file typedefs.hpp.

6.6 EXISTS Namespace Reference

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

Variables

```
• const int BOTH = -1
```

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

6.6.1 Detailed Description

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

6.6.2 Variable Documentation

6.6.2.1 AS_ONE

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

Definition at line 38 of file typedefs.hpp.

6.6.2.2 AS_ZERO

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

Definition at line 37 of file typedefs.hpp.

6.6.2.3 BOTH

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

Definition at line 31 of file typedefs.hpp.

6.6.2.4 NONE

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

Definition at line 32 of file typedefs.hpp.

6.6.2.5 ONE

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

Definition at line 33 of file typedefs.hpp.

6.6.2.6 TWO

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

Definition at line 34 of file typedefs.hpp.

6.6.2.7 UKNOWN

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

Definition at line 36 of file typedefs.hpp.

Chapter 7

Class Documentation

7.1 BArray < Cell_Type, Data_Type > Class Template Reference

Baseline class for binary arrays.

#include <barray-bones.hpp>

Public Member Functions

- bool operator== (const BArray< Cell_Type, Data_Type > &Array_)
- ∼BArray ()
- void out_of_range (uint i, uint j) const
- Cell_Type get_cell (uint i, uint j, bool check_bounds=true) const
- std::vector< Cell_Type > get_col_vec (uint i, bool check_bounds=true) const
- std::vector< Cell_Type > get_row_vec (uint i, bool check_bounds=true) const
- $\bullet \ \ \text{void} \ \ \underline{\text{get_col_vec}} \ \ (\text{std::vector} < \ \ \text{Cell_Type} > *x, \ \underline{\text{uint i, bool check_bounds=true}}) \ \ \text{const}$
- void get_row_vec (std::vector< Cell_Type > *x, uint i, bool check_bounds=true) const
- const Row_type< Cell_Type > & row (uint i, bool check_bounds=true) const
- const Col_type< Cell_Type > & col (uint i, bool check_bounds=true) const
- Entries < Cell_Type > get_entries () const

Get the edgelist.

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

Constructors

Parameters

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

• BArray ()

Zero-size array.

BArray (uint N_, uint M_)

Empty array.

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

Edgelist with data.

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

Edgelist with no data (simpler)

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

Move operator.

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

Queries

is_empty queries a single cell. nrow, ncol, and nnozero return the number of rows, columns, and non-zero cells respectively.

Parameters

i,j	Coordinates
check_bounds	If false avoids checking bounds.

- bool is empty (uint i, uint j, bool check bounds=true) const
- uint nrow () const noexcept
- uint ncol () const noexcept
- uint nnozero () const noexcept
- Cell< Cell_Type > default_val () const

Cell-wise insertion/deletion

Parameters

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

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

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

Column/row wise interchange

- void swap_rows (uint i0, uint i1, bool check_bounds=true)
- void swap_cols (uint j0, uint j1, bool check_bounds=true)
- void zero row (uint i, bool check bounds=true)
- void zero_col (uint j, bool check_bounds=true)

Arithmetic operators

```
• BArray< Cell_Type, Data_Type > & operator+= (const BArray< Cell_Type, Data_Type > &rhs)
```

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

Public Attributes

bool visited = false

Friends

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

7.1.1 Detailed Description

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

Baseline class for binary arrays.

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

Template Parameters

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

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

7.1.2 Constructor & Destructor Documentation

7.1.2.1 BArray() [1/6]

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

Zero-size array.

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

7.1.2.2 BArray() [2/6]

Empty array.

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

7.1.2.3 BArray() [3/6]

Edgelist with data.

7.1.2.4 BArray() [4/6]

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

Edgelist with no data (simpler)

7.1.2.5 BArray() [5/6]

Copy constructor.

7.1.2.6 BArray() [6/6]

Move operator.

7.1.2.7 \sim BArray()

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

7.1.3 Member Function Documentation

7.1.3.1 clear()

7.1.3.2 col()

7.1.3.3 D() [1/2]

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

7.1.3.4 D() [2/2]

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

7.1.3.5 default_val()

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

7.1.3.6 get_cell()

7.1.3.7 get_col_vec() [1/2]

7.1.3.8 get_col_vec() [2/2]

7.1.3.9 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>

7.1.3.10 get_row_vec() [1/2]

7.1.3.11 get_row_vec() [2/2]

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

7.1.3.12 insert_cell() [1/3]

7.1.3.13 insert_cell() [2/3]

7.1.3.14 insert_cell() [3/3]

7.1.3.15 is_empty()

7.1.3.16 ncol()

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

7.1.3.17 nnozero()

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

7.1.3.18 nrow()

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

7.1.3.19 operator()() [1/2]

7.1.3.20 operator()() [2/2]

7.1.3.21 operator*=()

7.1.3.22 operator+=() [1/3]

7.1.3.23 operator+=() [2/3]

7.1.3.24 operator+=() [3/3]

7.1.3.25 operator-=() [1/3]

7.1.3.26 operator-=() [2/3]

7.1.3.27 operator-=() [3/3]

7.1.3.28 operator/=()

7.1.3.29 operator=() [1/2]

Move assignment.

7.1.3.30 operator=() [2/2]

Assignment constructor.

7.1.3.31 operator==()

7.1.3.32 out of range()

7.1.3.33 print()

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

7.1.3.34 reserve()

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

7.1.3.35 resize()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
void BArray< Cell_Type, Data_Type >::resize (
          uint N_,
          uint M_ )
```

7.1.3.36 rm_cell()

7.1.3.37 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
```

7.1.3.38 set data()

Set the data object.

Parameters

```
data_
delete_←
data_
```

7.1.3.39 swap_cells()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
void BArray< Cell_Type, Data_Type >::swap_cells (
    uint i0,
    uint j0,
    uint i1,
    uint j1,
    bool check_bounds = true,
    int check_exists = CHECK::BOTH,
    int * report = nullptr )
```

7.1.3.40 swap_cols()

7.1.3.41 swap_rows()

7.1.3.42 toggle cell()

7.1.3.43 toggle_lock()

7.1.3.44 transpose()

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

7.1.3.45 zero_col()

7.1.3.46 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
```

7.2.1 Detailed Description

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

• bool operator== (const Cell_Type &val) const

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 BArrayDense< Cell_Type, Data_Type > Class Template Reference

Baseline class for binary arrays.

#include <barraydense-bones.hpp>

Public Member Functions

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

Get the edgelist.

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

Constructors

Parameters

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

• BArrayDense ()

Zero-size array.

• BArrayDense (uint N_, uint M_)

Emptv arrav.

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

Edgelist with data.

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

Edgelist with no data (simpler)

- BArrayDense (const BArrayDense < Cell_Type, Data_Type > &Array_, bool copy_data=false)
 Copy constructor.
- BArrayDense Cell_Type, Data_Type > & operator= (const BArrayDense Cell_Type, Data_Type > & Array_)

Assignment constructor.

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

Move operator.

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

Move assignment.

- void set_data (Data_Type *data_, bool delete_data_=false)
 Set the data object.
- Data_Type * D ()
- const Data_Type * D () const

Queries

is_empty queries a single cell. nrow, ncol, and nnozero return the number of rows, columns, and non-zero cells respectively.

Parameters

i,j	Coordinates
check_bounds	If false avoids checking bounds.

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

Cell-wise insertion/deletion

Parameters

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

- BArrayDense< Cell_Type, Data_Type > & operator+= (const std::pair< uint, uint > &coords)
- BArrayDense< Cell_Type, Data_Type > & operator-= (const std::pair< uint, uint > &coords)
- BArrayCell< Cell_Type, Data_Type > operator() (uint i, uint j, bool check_bounds=true)
- const BArrayCell_const< Cell_Type, Data_Type > operator() (uint i, uint j, bool check_bounds=true) const
- void rm_cell (uint i, uint j, bool check_bounds=true, bool check_exists=true)
- void insert_cell (uint i, uint 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

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

Public Attributes

bool visited = false

Friends

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

7.4.1 Detailed Description

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

Baseline class for binary arrays.

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

Template Parameters

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

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

7.4.2 Constructor & Destructor Documentation

7.4.2.1 BArrayDense() [1/6]

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

Zero-size array.

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

7.4.2.2 BArrayDense() [2/6]

Empty array.

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

7.4.2.3 BArrayDense() [3/6]

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

Edgelist with data.

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

7.4.2.4 BArrayDense() [4/6]

Edgelist with no data (simpler)

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

7.4.2.5 BArrayDense() [5/6]

Copy constructor.

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

7.4.2.6 BArrayDense() [6/6]

Move operator.

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

7.4.2.7 ∼BArrayDense()

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

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

7.4.3 Member Function Documentation

7.4.3.1 clear()

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

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

7.4.3.2 col()

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

7.4.3.3 D() [1/2]

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

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

7.4.3.4 D() [2/2]

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

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

7.4.3.5 default_val()

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

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

7.4.3.6 get_cell()

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

7.4.3.7 get_col_vec() [1/2]

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

7.4.3.8 get_col_vec() [2/2]

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

7.4.3.9 get_entries()

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

Get the edgelist.

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

Returns

```
Entries<Cell_Type>
```

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

7.4.3.10 get row_vec() [1/2]

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

7.4.3.11 get_row_vec() [2/2]

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

7.4.3.12 insert_cell() [1/3]

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

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

7.4.3.13 insert_cell() [2/3]

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

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

7.4.3.14 insert_cell() [3/3]

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

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

7.4.3.15 is_empty()

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

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

7.4.3.16 ncol()

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

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

7.4.3.17 nnozero()

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

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

7.4.3.18 nrow()

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

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

7.4.3.19 operator()() [1/2]

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

7.4.3.20 operator()() [2/2]

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

7.4.3.21 operator*=()

7.4.3.22 operator+=() [1/3]

7.4.3.23 operator+=() [2/3]

7.4.3.24 operator+=() [3/3]

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

7.4.3.25 operator-=() [1/3]

7.4.3.26 operator-=() [2/3]

7.4.3.27 operator-=() [3/3]

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

7.4.3.28 operator/=()

7.4.3.29 operator=() [1/2]

Move assignment.

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

7.4.3.30 operator=() [2/2]

Assignment constructor.

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

7.4.3.31 operator==()

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

7.4.3.32 out_of_range()

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

7.4.3.33 print()

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

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

7.4.3.34 reserve()

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

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

7.4.3.35 resize()

```
template<typename Cell_Type , typename Data_Type >
void BArrayDense< Cell_Type, Data_Type >::resize (
          uint N_,
          uint M_ ) [inline]
```

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

7.4.3.36 rm_cell()

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

7.4.3.37 row()

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

7.4.3.38 set_data()

Set the data object.

Parameters

data_	
delete_←	
data_	

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

7.4.3.39 swap_cells()

```
template<typename Cell_Type , typename Data_Type >
void BArrayDense< Cell_Type, Data_Type >::swap_cells (
          uint i0,
          uint j0,
          uint i1,
          uint j1,
          bool check_bounds = true,
          int check_exists = CHECK::BOTH,
          int * report = nullptr ) [inline]
```

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

7.4.3.40 swap_cols()

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

7.4.3.41 swap_rows()

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

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

7.4.3.42 toggle_cell()

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

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

7.4.3.43 toggle_lock()

7.4.3.44 transpose()

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

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

7.4.3.45 zero_col()

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

7.4.3.46 zero_row()

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

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

7.4.4 Friends And Related Function Documentation

7.4.4.1 BArrayCell < Cell_Type, Data_Type >

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

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

7.4.4.2 BArrayCell_const< Cell_Type, Data_Type >

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

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

7.4.5 Member Data Documentation

7.4.5.1 visited

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

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

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

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

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

7.5 BArrayDenseCell< Cell_Type, Data_Type > Class Template Reference

#include <barraydensecell-bones.hpp>

Public Member Functions

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

7.5.1 Detailed Description

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

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

7.5.2 Constructor & Destructor Documentation

7.5.2.1 BArrayDenseCell()

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

7.5.2.2 ~BArrayDenseCell()

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

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

7.5.3 Member Function Documentation

7.5.3.1 operator Cell_Type()

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

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

7.5.3.2 operator*=()

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

7.5.3.3 operator+=()

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

7.5.3.4 operator-=()

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

7.5.3.5 operator/=()

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

7.5.3.6 operator=()

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

7.5.3.7 operator==()

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

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

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

7.6 BArrayDenseCell_const< Cell_Type, Data_Type > Class Template Reference

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

Public Member Functions

- BArrayDenseCell_const (const BArrayDense < Cell_Type, Data_Type > *Array_, uint i_, uint j_, bool check
 _bounds=true)
- ∼BArrayDenseCell const ()
- · operator Cell Type () const
- bool operator== (const Cell_Type &val) const
- bool operator!= (const Cell_Type &val) const
- bool operator< (const Cell_Type &val) const
- bool operator> (const Cell_Type &val) const
- bool operator<= (const Cell_Type &val) const
- bool operator>= (const Cell_Type &val) const

7.6.1 Detailed Description

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

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

7.6.2 Constructor & Destructor Documentation

7.6.2.1 BArrayDenseCell_const()

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

7.6.2.2 ~BArrayDenseCell const()

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

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

7.6.3 Member Function Documentation

7.6.3.1 operator Cell_Type()

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

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

7.6.3.2 operator"!=()

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

7.6.3.3 operator<()

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

7.6.3.4 operator<=()

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

7.6.3.5 operator==()

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

7.6.3.6 operator>()

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

7.6.3.7 operator>=()

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

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

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

7.7 BArrayVector < Cell_Type, Data_Type > Class Template Reference

Row or column of a BArray

#include <barrayvector-bones.hpp>

Public Member Functions

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

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

7.7.1 Detailed Description

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

Row or column of a BArray

Template Parameters

Cell_Type	
Data_Type	

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

7.7.2 Constructor & Destructor Documentation

7.7.2.1 BArrayVector()

Construct a new BArrayVector object.

Parameters

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

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

7.7.2.2 ∼BArrayVector()

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

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

7.7.3 Member Function Documentation

7.7.3.1 begin()

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

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

7.7.3.2 end()

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

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

7.7.3.3 is_col()

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

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

7.7.3.4 is_row()

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

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

7.7.3.5 operator std::vector< Cell_Type >()

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

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

7.7.3.6 operator*=()

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

7.7.3.7 operator+=()

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

7.7.3.8 operator-=()

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

7.7.3.9 operator/=()

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

7.7.3.10 operator=()

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

7.7.3.11 operator==()

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

7.7.3.12 size()

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

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

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

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

7.8 BArrayVector_const< Cell_Type, Data_Type > Class Template Reference

#include <barrayvector-bones.hpp>

Public Member Functions

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

7.8.1 Detailed Description

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

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

7.8.2 Constructor & Destructor Documentation

7.8.2.1 BArrayVector_const()

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

7.8.2.2 ~BArrayVector_const()

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

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

7.8.3 Member Function Documentation

7.8.3.1 begin()

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

7.8.3.2 end()

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

7.8.3.3 is_col()

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

7.8.3.4 is_row()

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

7.8.3.5 operator std::vector< Cell_Type >()

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

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

7.8.3.6 operator"!=()

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

7.8.3.7 operator<()

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

7.8.3.8 operator<=()

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

7.8.3.9 operator==()

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

7.8.3.10 operator>()

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

7.8.3.11 operator>=()

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

7.8.3.12 size()

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

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

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

7.9 Cell < Cell Type > Class Template Reference

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

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

Public Member Functions

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

Public Attributes

- Cell_Type value
- bool visited

7.9.1 Detailed Description

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

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

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

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

7.9.2 Constructor & Destructor Documentation

7.9.2.1 Cell() [1/7]

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

7.9.2.2 Cell() [2/7]

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

7.9.2.3 ∼CeII()

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

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

7.9.2.4 Cell() [3/7]

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

7.9.2.5 Cell() [4/7]

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

7.9.2.6 Cell() [5/7]

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

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

7.9.2.7 Cell() [6/7]

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

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

7.9.2.8 Cell() [7/7]

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

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

7.9.3 Member Function Documentation

7.9.3.1 add() [1/4]

7.9.3.2 add() [2/4]

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

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

7.9.3.3 add() [3/4]

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

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

7.9.3.4 add() [4/4]

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

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

7.9.3.5 operator Cell_Type()

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

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

7.9.3.6 operator"!=()

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

7.9.3.7 operator=() [1/2]

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

7.9.3.8 operator=() [2/2]

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

7.9.3.9 operator==()

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

7.9.4 Member Data Documentation

7.9.4.1 value

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

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

7.9.4.2 visited

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

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

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

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

7.10 ConstBArrayRowlter< Cell_Type, Data_Type > Class Template Reference

#include <barray-iterator.hpp>

Collaboration diagram for ConstBArrayRowlter< Cell_Type, Data_Type >:



Public Member Functions

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

Public Attributes

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

7.10.1 Detailed Description

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

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

7.10.2 Constructor & Destructor Documentation

7.10.2.1 ConstBArrayRowlter()

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

7.10.2.2 ~ConstBArrayRowlter()

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

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

7.10.3 Member Data Documentation

7.10.3.1 Array

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

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

7.10.3.2 current_col

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

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

7.10.3.3 current_row

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

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

7.10.3.4 iter

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

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

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

include/barry/barray-iterator.hpp

7.11 Counter< Array_Type, Data_Type > Class Template Reference

A counter function based on change statistics.

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

Public Member Functions

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

Creator passing a counter and an initializer

Parameters

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

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

Move assignment.

Public Attributes

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

7.11.1 Detailed Description

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

A counter function based on change statistics.

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

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

7.11.2 Constructor & Destructor Documentation

7.11.2.1 Counter() [1/4]

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

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

7.11.2.2 Counter() [2/4]

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

7.11.2.3 Counter() [3/4]

Copy constructor.

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

7.11.2.4 Counter() [4/4]

Move constructor.

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

7.11.2.5 ~Counter()

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

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

7.11.3 Member Function Documentation

7.11.3.1 count()

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

7.11.3.2 init()

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

7.11.3.3 operator=() [1/2]

Copy assignment.

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

7.11.3.4 operator=() [2/2]

Move assignment.

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

7.11.4 Member Data Documentation

7.11.4.1 count_fun

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

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

7.11.4.2 data

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

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

7.11.4.3 delete_data

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

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

7.11.4.4 desc

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

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

7.11.4.5 init_fun

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

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

7.11.4.6 name

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

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

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

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

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

Vector of counters.

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

Public Member Functions

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

Copy constructor.

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

Move constructor.

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

Move assignment constructor.

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

Returns a pointer to a particular counter.

• std::size_t size () const noexcept

Number of counters in the set.

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

7.12.1 Detailed Description

Vector of counters.

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

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

7.12.2 Constructor & Destructor Documentation

7.12.2.1 Counters() [1/3]

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

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

7.12.2.2 ∼Counters()

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

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

7.12.2.3 Counters() [2/3]

Copy constructor.

Parameters



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

7.12.2.4 Counters() [3/3]

Move constructor.

Parameters



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

7.12.3 Member Function Documentation

7.12.3.1 add counter() [1/3]

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

7.12.3.2 add_counter() [2/3]

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

7.12.3.3 add_counter() [3/3]

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

7.12.3.4 clear()

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

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

7.12.3.5 operator=() [1/2]

Copy assignment constructor.

Parameters

counter←	
_	

Returns

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

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

7.12.3.6 operator=() [2/2]

Move assignment constructor.

Parameters



Returns

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

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

7.12.3.7 operator[]()

Returns a pointer to a particular counter.

Parameters

```
idx Id of the counter
```

Returns

Counter<Array_Type,Data_Type>*

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

7.12.3.8 size()

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

Number of counters in the set.

Returns

uint

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

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

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

7.13 Entries < Cell_Type > Class Template Reference

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

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

Public Member Functions

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

Public Attributes

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

7.13.1 Detailed Description

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

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

Template Parameters

```
Cell_Type Any type
```

Definition at line 59 of file typedefs.hpp.

7.13.2 Constructor & Destructor Documentation

7.13.2.1 Entries() [1/2]

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

Definition at line 65 of file typedefs.hpp.

7.13.2.2 Entries() [2/2]

Definition at line 66 of file typedefs.hpp.

7.13.2.3 ∼Entries()

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

Definition at line 73 of file typedefs.hpp.

7.13.3 Member Function Documentation

7.13.3.1 resize()

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

Definition at line 75 of file typedefs.hpp.

7.13.4 Member Data Documentation

7.13.4.1 source

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

Definition at line 61 of file typedefs.hpp.

7.13.4.2 target

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

Definition at line 62 of file typedefs.hpp.

7.13.4.3 val

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

Definition at line 63 of file typedefs.hpp.

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

• include/barry/typedefs.hpp

7.14 Flock Class Reference

A Flock is a group of Geese.

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

7.14 Flock Class Reference 93

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 * get_counters ()
- phylocounters::PhyloSupport * get support ()

Returns the joint likelihood of the model.

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

Access the i-th geese element.

Information about the model

- unsigned int nfuns () const noexcept
- unsigned int ntrees () const noexcept
- std::vector< unsigned int > nnodes () const noexcept
- std::vector< unsigned int > nleafs () const noexcept
- unsigned int nterms () const
- · unsigned int support size () const noexcept

Public Attributes

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

7.14.1 Detailed Description

A Flock is a group of Geese.

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

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

7.14.2 Constructor & Destructor Documentation

7.14.2.1 Flock()

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

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

7.14.2.2 ∼Flock()

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

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

7.14.3 Member Function Documentation

7.14.3.1 add_data()

Add a tree to the flock.

Parameters

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

Returns

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

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

7.14.3.2 get_counters()

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

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

7.14 Flock Class Reference 95

7.14.3.3 get_support()

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

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

7.14.3.4 init()

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

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

7.14.3.5 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	ence When true (default) will compute the likelihood using the reduced sequence,	
	which is faster.	

Returns

double

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

7.14.3.6 nfuns()

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

Definition at line 107 of file flock-meet.hpp.

7.14.3.7 nleafs()

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

Definition at line 130 of file flock-meet.hpp.

7.14.3.8 nnodes()

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

Definition at line 119 of file flock-meet.hpp.

7.14.3.9 nterms()

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

Definition at line 142 of file flock-meet.hpp.

7.14.3.10 ntrees()

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

Definition at line 113 of file flock-meet.hpp.

7.14.3.11 operator()()

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

Access the i-th geese element.

Parameters

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

Returns

Geese*

7.14 Flock Class Reference 97

Definition at line 155 of file flock-meet.hpp.

7.14.3.12 set_seed()

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

7.14.3.13 support_size()

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

Definition at line 149 of file flock-meet.hpp.

7.14.4 Member Data Documentation

7.14.4.1 dat

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

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

7.14.4.2 initialized

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

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

7.14.4.3 model

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

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

7.14.4.4 nfunctions

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

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

7.14.4.5 rengine

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

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

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

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

7.15 FreqTable < T > Class Template Reference

Database of statistics.

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

Public Member Functions

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

7.15.1 Detailed Description

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

Database of statistics.

This is mostly used in Support.

Definition at line 16 of file statsdb.hpp.

7.15.2 Constructor & Destructor Documentation

7.15.2.1 FreqTable()

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

Definition at line 28 of file statsdb.hpp.

7.15.2.2 ∼FreqTable()

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

Definition at line 29 of file statsdb.hpp.

7.15.3 Member Function Documentation

7.15.3.1 add()

Definition at line 47 of file statsdb.hpp.

7.15.3.2 as_vector()

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

Definition at line 61 of file statsdb.hpp.

7.15.3.3 clear()

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

Definition at line 83 of file statsdb.hpp.

7.15.3.4 get_data()

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

Definition at line 73 of file statsdb.hpp.

7.15.3.5 get_data_ptr()

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

Definition at line 78 of file statsdb.hpp.

7.15.3.6 print()

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

Definition at line 102 of file statsdb.hpp.

7.15.3.7 reserve()

Definition at line 89 of file statsdb.hpp.

7.15.3.8 size()

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

Definition at line 126 of file statsdb.hpp.

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

• include/barry/statsdb.hpp

7.16 Geese Class Reference

Annotated Phylo Model.

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

Public Member Functions

- ~Geese ()
- void init ()
- void inherit_support (const Geese &model_, bool delete_support_=false)
- void calc sequence (Node *n=nullptr)
- void calc reduced sequence ()
- double likelihood (const std::vector< double > &par, bool as_log=false, bool use_reduced_sequence=true)
- double likelihood_exhaust (const std::vector< double > &par)
- std::vector< double > get_probabilities () const
- void set_seed (const unsigned int &s)
- std::vector< std::vector< unsigned int > > simulate (const std::vector< double > &par)
- std::vector< std::vector< double >> observed counts ()
- void print_observed_counts ()
- void init_node (Node &n)
- void update_annotations (unsigned int nodeid, std::vector< unsigned int > newann)
- std::mt19937 * get_rengine ()
- phylocounters::PhyloCounters * get_counters ()
- phylocounters::PhyloModel * get model ()
- phylocounters::PhyloSupport * get_support ()
- std::vector< std::vector< bool > > get_states ()

Construct a new Geese object

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

Parameters

annotations	A vector of vectors with annotations. It should be of length k (number of functions). Each vector should be of length N (equal to the number of nodes, including interior). Possible values are 0, 1, and 9.
geneid	Id of the gene. It should be of length ${\tt N}.$
parent	Id of the parent gene. Also of length ${\tt N}$

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

Information about the model

Parameters

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

• unsigned int nfuns () const noexcept

Number of functions analyzed.

· unsigned int nnodes () const noexcept

Number of nodes (interior + leaf)

• unsigned int nleafs () const noexcept

Number of leaf.

• unsigned int nterms () const

Number of terms included.

• unsigned int support_size () const noexcept

Number of unique sets of sufficient stats.

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

Check polytomies and return the largest.

Geese prediction

Calculate the conditional probability

Parameters

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

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

Returns

std::vector< double > Returns the posterior probability

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

Public Attributes

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

7.16.1 Detailed Description

Annotated Phylo Model.

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

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

7.16.2 Constructor & Destructor Documentation

7.16.2.1 Geese() [1/4]

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

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

7.16.2.2 Geese() [2/4]

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

7.16.2.3 Geese() [3/4]

Definition at line 163 of file geese-meat-constructors.hpp.

7.16.2.4 Geese() [4/4]

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

7.16.2.5 ∼Geese()

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

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

7.16.3 Member Function Documentation

7.16.3.1 calc_reduced_sequence()

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

Definition at line 225 of file geese-meat.hpp.

7.16.3.2 calc_sequence()

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

Definition at line 182 of file geese-meat.hpp.

7.16.3.3 get_counters()

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

Definition at line 456 of file geese-meat.hpp.

7.16.3.4 get_model()

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

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

7.16.3.5 get_probabilities()

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

Definition at line 268 of file geese-meat.hpp.

7.16.3.6 get_rengine()

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

Definition at line 452 of file geese-meat.hpp.

7.16.3.7 get_states()

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

Definition at line 468 of file geese-meat.hpp.

7.16.3.8 get_support()

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

Definition at line 464 of file geese-meat.hpp.

7.16.3.9 inherit_support()

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

7.16.3.10 init()

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

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

7.16.3.11 init_node()

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

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

7.16.3.12 likelihood()

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

7.16.3.13 likelihood_exhaust()

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

7.16 Geese Class Reference 107

7.16.3.14 nfuns()

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

Number of functions analyzed.

Definition at line 284 of file geese-meat.hpp.

7.16.3.15 nleafs()

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

Number of leaf.

Definition at line 292 of file geese-meat.hpp.

7.16.3.16 nnodes()

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

Number of nodes (interior + leaf)

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

7.16.3.17 nterms()

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

Number of terms included.

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

7.16.3.18 observed_counts()

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

Definition at line 342 of file geese-meat.hpp.

7.16.3.19 operator=() [1/2]

7.16.3.20 operator=() [2/2]

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

7.16.3.21 parse_polytomies()

Check polytomies and return the largest.

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

7.16.3.22 predict()

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

7.16.3.23 predict_backend()

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

7.16 Geese Class Reference 109

7.16.3.24 print_observed_counts()

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

Definition at line 390 of file geese-meat.hpp.

7.16.3.25 set_seed()

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

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

7.16.3.26 simulate()

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

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

7.16.3.27 support_size()

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

Number of unique sets of sufficient stats.

Definition at line 309 of file geese-meat.hpp.

7.16.3.28 update_annotations()

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

Definition at line 159 of file geese-meat.hpp.

7.16.4 Member Data Documentation

7.16.4.1 delete_rengine

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

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

7.16.4.2 delete_support

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

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

7.16.4.3 initialized

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

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

7.16.4.4 map_to_nodes

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

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

7.16.4.5 nfunctions

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

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

7.16.4.6 nodes

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

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

7.16.4.7 reduced_sequence

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

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

7.16.4.8 sequence

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

Definition at line 100 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

7.17 Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data Rule Dyn Type > Class Template Reference

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

#include <model-bones.hpp>

Public Member Functions

- void set rengine (std::mt19937 *rengine , bool delete =false)
- void set_seed (unsigned int s)
- Model ()
- Model (uint size_)
- Model (const Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > &Model ←
 _)
- Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > & operator= (const Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > &Model_)
- \sim 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)

- const std::mt19937 * get rengine () const
- Counters < Array_Type, Data_Counter_Type > * get_counters ()
- Rules< Array_Type, Data_Rule_Type > * get_rules ()
- Rules< Array_Type, Data_Rule_Dyn_Type > * get_rules_dyn ()
- Support < Array Type, Data Counter Type, Data Rule Type, Data Rule Dyn Type > * get support ()

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

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

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

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

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

- void add_rule (Rule < Array_Type, Data_Rule_Type > &rule)
- void add_rule (Rule < Array_Type, Data_Rule_Type > *rule)
- void add_rule (Rule_fun_type< Array_Type, Data_Rule_Type > count_fun_, Data_Rule_Type *data_←
 =nullptr, bool delete data =false)
- void set_rules (Rules < Array_Type, Data_Rule_Type > *rules_)
- void add_rule_dyn (Rule < Array_Type, Data_Rule_Dyn_Type > &rule)
- void add_rule_dyn (Rule < Array_Type, Data_Rule_Dyn_Type > *rule)
- void set_rules_dyn (Rules < Array_Type, Data_Rule_Dyn_Type > *rules_)

Likelihood functions.

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

Parameters

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

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

Extract elements by index

Parameters

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

- double get_norm_const (const std::vector< double > ¶ms, const uint &i, bool as_log=false)
- const std::vector< Array_Type > * get_pset (const uint &i)
- const std::vector< std::vector< double > > * get stats (const uint &i)

Size of the model

Number of different supports included in the model

This will return the size of stats.

Returns

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

- unsigned int size () const noexcept
- unsigned int size unique () const noexcept
- · unsigned int nterms () const noexcept
- · unsigned int support size () const noexcept

7.17.1 Detailed Description

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

class Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >

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

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

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

Template Parameters

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

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

7.17.2 Constructor & Destructor Documentation

7.17.2.1 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.

7.17.2.2 Model() [2/3]

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

7.17.2.3 Model() [3/3]

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

7.17.2.4 ~Model()

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

Definition at line 204 of file model-bones.hpp.

7.17.3 Member Function Documentation

7.17.3.1 add_array()

Adds an array to the support of not already included.

Parameters

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

Returns

The number of the array.

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

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

7.17.3.2 add_counter() [1/3]

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

7.17.3.3 add_counter() [2/3]

Definition at line 176 of file model-meat.hpp.

7.17.3.4 add_counter() [3/3]

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

7.17.3.5 add_rule() [1/3]

Definition at line 225 of file model-meat.hpp.

7.17.3.6 add_rule() [2/3]

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

7.17.3.7 add_rule() [3/3]

Definition at line 244 of file model-meat.hpp.

7.17.3.8 add_rule_dyn() [1/3]

Definition at line 279 of file model-meat.hpp.

7.17.3.9 add_rule_dyn() [2/3]

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

7.17.3.10 add_rule_dyn() [3/3]

Definition at line 298 of file model-meat.hpp.

7.17.3.11 get_counters()

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

Counters< Array_Type, Data_Counter_Type > * Model< Array_Type, Data_Counter_Type, Data_Rule

_Type, Data_Rule_Dyn_Type >::get_counters [inline]
```

Definition at line 713 of file model-meat.hpp.

7.17.3.12 get_norm_const()

Definition at line 567 of file model-meat.hpp.

7.17.3.13 get pset()

Definition at line 599 of file model-meat.hpp.

7.17.3.14 get_rengine()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
const std::mt19937 * Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type
>::get_rengine [inline]
```

Definition at line 707 of file model-meat.hpp.

7.17.3.15 get_rules()

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

Definition at line 719 of file model-meat.hpp.

7.17.3.16 get_rules_dyn()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
Rules< Array_Type, Data_Rule_Dyn_Type > * Model< Array_Type, Data_Counter_Type, Data_Rule_\top Type, Data_Rule_Dyn_Type >::get_rules_dyn [inline]
```

Definition at line 725 of file model-meat.hpp.

7.17.3.17 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 612 of file model-meat.hpp.

7.17.3.18 get_support()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > * Model< Array

_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::get_support [inline]
```

Definition at line 731 of file model-meat.hpp.

7.17.3.19 likelihood() [1/3]

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

7.17.3.20 likelihood() [2/3]

Definition at line 493 of file model-meat.hpp.

7.17.3.21 likelihood() [3/3]

Definition at line 420 of file model-meat.hpp.

7.17.3.22 likelihood_total()

Definition at line 527 of file model-meat.hpp.

7.17.3.23 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 653 of file model-meat.hpp.

7.17.3.24 operator=()

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

7.17.3.25 print_stats()

Definition at line 624 of file model-meat.hpp.

7.17.3.26 sample() [1/2]

7.17.3.27 sample() [2/2]

Definition at line 671 of file model-meat.hpp.

7.17.3.28 set_counters()

Definition at line 205 of file model-meat.hpp.

7.17.3.29 set_keygen()

Definition at line 159 of file model-meat.hpp.

7.17.3.30 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 176 of file model-bones.hpp.

7.17.3.31 set_rules()

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

7.17.3.32 set rules dyn()

Definition at line 315 of file model-meat.hpp.

7.17.3.33 set_seed()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarrow 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 186 of file model-bones.hpp.

7.17.3.34 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 643 of file model-meat.hpp.

7.17.3.35 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 648 of file model-meat.hpp.

7.17.3.36 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 151 of file model-meat.hpp.

7.17.3.37 support_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 >::support_size
[inline], [noexcept]
```

Definition at line 660 of file model-meat.hpp.

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

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

7.18 NetCounterData Class Reference

Data class used to store arbitrary uint or double vectors.

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

Public Member Functions

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

Public Attributes

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

7.18.1 Detailed Description

Data class used to store arbitrary uint or double vectors.

Definition at line 61 of file network.hpp.

7.18.2 Constructor & Destructor Documentation

7.18.2.1 NetCounterData() [1/2]

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

Definition at line 67 of file network.hpp.

7.18.2.2 NetCounterData() [2/2]

Definition at line 68 of file network.hpp.

7.18.2.3 ∼NetCounterData()

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

Definition at line 73 of file network.hpp.

7.18.3 Member Data Documentation

7.18.3.1 indices

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

Definition at line 64 of file network.hpp.

7.18.3.2 numbers

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

Definition at line 65 of file network.hpp.

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

• include/barry/counters/network.hpp

7.19 NetworkData Class Reference

Data class for Networks.

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

Public Member Functions

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

Constructor using a single attribute.

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

Constructor using multiple attributes.

• \sim NetworkData ()

Public Attributes

- bool directed = true
- std::vector< std::vector< double >> vertex_attr

7.19.1 Detailed Description

Data class for Networks.

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

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

Definition at line 24 of file network.hpp.

7.19.2 Constructor & Destructor Documentation

7.19.2.1 NetworkData() [1/3]

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

Definition at line 30 of file network.hpp.

7.19.2.2 NetworkData() [2/3]

Constructor using a single attribute.

Parameters

vert	tex_←	Double vector of length equal to the number of vertices in the data.
attr_	_	
dire	ected_	When true the graph as treated as directed.

Definition at line 38 of file network.hpp.

7.19.2.3 NetworkData() [3/3]

Constructor using multiple attributes.

Parameters

vertex_← attr_	Vector of double vectors. The size equals to the number of attributes to be created. Each 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.

7.20 Node Class Reference 127

7.19.2.4 ~NetworkData()

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

Definition at line 56 of file network.hpp.

7.19.3 Member Data Documentation

7.19.3.1 directed

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

Definition at line 27 of file network.hpp.

7.19.3.2 vertex_attr

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

Definition at line 28 of file network.hpp.

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

• include/barry/counters/network.hpp

7.20 Node Class Reference

A single node for the model.

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

Collaboration diagram for Node:



Public Member Functions

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

Construct a new Node object

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

Public Attributes

· unsigned int id

Id of the node (as specified in the input)

· unsigned int ord

Order in which the node was created.

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

Observed annotations (only defined for Geese)

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

Arrays given all possible states.

Node * parent = nullptr

Parent node.

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

Offspring nodes.

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

ID of the array in the model.

- bool visited = false
- std::vector< double > subtree_prob

Induced subtree probabilities.

std::vector< double > probability

The probability of observing each state.

7.20.1 Detailed Description

A single node for the model.

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

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

7.20.2 Constructor & Destructor Documentation

7.20 Node Class Reference 129

7.20.2.1 Node() [1/5]

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

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

7.20.2.2 Node() [2/5]

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

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

7.20.2.3 Node() [3/5]

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

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

7.20.2.4 Node() [4/5]

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

7.20.2.5 Node() [5/5]

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

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

7.20.2.6 ∼Node()

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

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

7.20.3 Member Function Documentation

7.20.3.1 get_parent()

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

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

7.20.3.2 is_leaf()

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

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

7.20.3.3 noffspring()

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

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

7.20.4 Member Data Documentation

7.20.4.1 annotations

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

Observed annotations (only defined for Geese)

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

7.20 Node Class Reference 131

7.20.4.2 array

phylocounters::PhyloArray Node::array

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

7.20.4.3 arrays

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

Arrays given all possible states.

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

7.20.4.4 duplication

bool Node::duplication

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

7.20.4.5 id

unsigned int Node::id

Id of the node (as specified in the input)

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

7.20.4.6 narray

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

ID of the array in the model.

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

7.20.4.7 offspring

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

Offspring nodes.

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

7.20.4.8 ord

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

Order in which the node was created.

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

7.20.4.9 parent

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

Parent node.

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

7.20.4.10 probability

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

The probability of observing each state.

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

7.20.4.11 subtree_prob

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

Induced subtree probabilities.

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

7.20.4.12 visited

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

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

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

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

7.21 NodeData Class Reference

Data definition for the PhyloArray class.

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

Public Member Functions

- NodeData ()
- 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

7.21.1 Detailed Description

Data definition for the PhyloArray class.

 $\textbf{Details about the available counters for $\tt PhyloArray objects can be found in the Phylo counters section. } \\$

This holds basic information about a given node.

Definition at line 23 of file phylo.hpp.

7.21.2 Constructor & Destructor Documentation

7.21.2.1 NodeData() [1/2]

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

Definition at line 41 of file phylo.hpp.

7.21.2.2 NodeData() [2/2]

Definition at line 43 of file phylo.hpp.

7.21.2.3 ~NodeData()

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

Definition at line 49 of file phylo.hpp.

7.21.3 Member Data Documentation

7.21.3.1 blengths

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

Branch length.

Definition at line 29 of file phylo.hpp.

7.21.3.2 duplication

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

Definition at line 39 of file phylo.hpp.

7.21.3.3 states

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

State of the parent node.

Definition at line 34 of file phylo.hpp.

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

• include/barry/counters/phylo.hpp

7.22 PhyloRuleDynData Class Reference

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

Public Member Functions

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

Public Attributes

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

7.22.1 Detailed Description

Definition at line 1189 of file phylo.hpp.

7.22.2 Constructor & Destructor Documentation

7.22.2.1 PhyloRuleDynData()

Definition at line 1196 of file phylo.hpp.

7.22.2.2 ~PhyloRuleDynData()

PhyloRuleDynData::~PhyloRuleDynData () [inline]

Definition at line 1205 of file phylo.hpp.

7.22.3 Member Data Documentation

7.22.3.1 counts

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

Definition at line 1191 of file phylo.hpp.

7.22.3.2 duplication

bool PhyloRuleDynData::duplication

Definition at line 1195 of file phylo.hpp.

7.22.3.3 lb

uint PhyloRuleDynData::lb

Definition at line 1193 of file phylo.hpp.

7.22.3.4 pos

uint PhyloRuleDynData::pos

Definition at line 1192 of file phylo.hpp.

7.22.3.5 ub

uint PhyloRuleDynData::ub

Definition at line 1194 of file phylo.hpp.

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

• include/barry/counters/phylo.hpp

7.23 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

7.23.1 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.

7.23.2 Constructor & Destructor Documentation

7.23.2.1 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.

7.23.2.2 PowerSet() [2/3]

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

7.23.2.3 PowerSet() [3/3]

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

7.23.2.4 ∼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.

7.23.3 Member Function Documentation

7.23.3.1 add_rule() [1/3]

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

7.23.3.2 add_rule() [2/3]

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

7.23.3.3 add_rule() [3/3]

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

7.23.3.4 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.

7.23.3.5 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.

7.23.3.6 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.

7.23.3.7 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.

7.23.3.8 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.

7.23.3.9 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.

7.23.3.10 operator[]()

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

7.23.3.11 reset()

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

7.23.3.12 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.

7.23.4 Member Data Documentation

7.23.4.1 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.

7.23.4.2 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.

7.23.4.3 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.

7.23.4.4 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.

7.23.4.5 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.

7.23.4.6 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.

7.23.4.7 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.

7.23.4.8 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

7.24 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)

7.24.1 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.

7.24.2 Constructor & Destructor Documentation

7.24.2.1 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.

7.24.2.2 Rule() [2/2]

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

7.24.2.3 ∼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.

7.24.3 Member Function Documentation

7.24.3.1 D()

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

Read/Write access to the data.

7.24.3.2 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

7.25 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)

7.25.1 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.

7.25.2 Constructor & Destructor Documentation

7.25.2.1 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.

7.25.2.2 Rules() [2/2]

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

7.25.2.3 ∼Rules()

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

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

7.25.3 Member Function Documentation

7.25.3.1 add_rule() [1/3]

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

7.25.3.2 add_rule() [2/3]

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

7.25.3.3 add_rule() [3/3]

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

7.25.3.4 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.

7.25.3.5 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.

7.25.3.6 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.

7.25.3.7 operator=()

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

7.25.3.8 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

7.26 StatsCounter< Array_Type, Data_Type > Class Template Reference

Count stats for a single Array.

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

Public Member Functions

StatsCounter (const Array Type *Array)

Creator of a StatsCounter

• StatsCounter ()

Can be created without setting the array.

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

Changes the reference array for the counting.

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

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

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

7.26.1 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.

7.26.2 Constructor & Destructor Documentation

7.26.2.1 StatsCounter() [1/2]

Creator of a StatsCounter

Parameters

Array←	A const pointer to a BArray.	
_		

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

7.26.2.2 StatsCounter() [2/2]

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

Can be created without setting the array.

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

7.26.2.3 ~StatsCounter()

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

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

7.26.3 Member Function Documentation

7.26.3.1 add_counter() [1/2]

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

7.26.3.2 add_counter() [2/2]

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

7.26.3.3 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.

7.26.3.4 count_current()

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

7.26.3.5 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.

7.26.3.6 get_counters()

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

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

7.26.3.7 reset_array()

Changes the reference array for the counting.

Parameters

Array←	A pointer to an array of class Array_Type.
_	

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

7.26.3.8 set_counters()

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

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

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

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

Compute the support of sufficient statistics.

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

Public Member Functions

Support (const Array_Type &Array_)

Constructor passing a reference Array.

Support (uint N_, uint M_)

Constructor specifying the dimensions of the array (empty).

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

Computes the entire support.

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

List current statistics.

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

Vector of couter functions.

Rules< Array_Type, Data_Rule_Type > * get_rules ()

Vector of static rules (cells to iterate).

Rules< Array_Type, Data_Rule_Dyn_Type > * get_rules_dyn ()

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

Resets the support calculator

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

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

- 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

7.27.1 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.

7.27.2 Constructor & Destructor Documentation

7.27.2.1 Support() [1/3]

Constructor passing a reference Array.

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

7.27.2.2 Support() [2/3]

Constructor specifying the dimensions of the array (empty).

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

7.27.2.3 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.

7.27.2.4 ~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.

7.27.3 Member Function Documentation

7.27.3.1 add_counter() [1/2]

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

7.27.3.2 add_counter() [2/2]

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

7.27.3.3 add_rule() [1/2]

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

7.27.3.4 add_rule() [2/2]

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

7.27.3.5 add_rule_dyn() [1/2]

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

7.27.3.6 add_rule_dyn() [2/2]

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

7.27.3.7 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.

7.27.3.8 get_counters()

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

Counters< Array_Type, Data_Counter_Type > * Support< Array_Type, Data_Counter_Type, Data_←

Rule_Type, Data_Rule_Dyn_Type >::get_counters [inline]
```

Vector of couter functions.

Definition at line 349 of file support-meat.hpp.

7.27.3.9 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.

7.27.3.10 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.

7.27.3.11 get_current_stats()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename Data_Rule_Dyn_Type >
std::vector< double > * Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Counter_Type >::get_current_stats [inline]
```

List current statistics.

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

7.27.3.12 get_data()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
const FreqTable & Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type
>::get_data [inline]
```

Definition at line 344 of file support-meat.hpp.

7.27.3.13 get_rules()

```
template<typename Array_Type , typename Data_Counter_Type , typename Data_Rule_Type , typename
Data_Rule_Dyn_Type >
Rules< Array_Type, Data_Rule_Type > * Support< Array_Type, Data_Counter_Type, Data_Rule_Type,
Data_Rule_Dyn_Type >::get_rules [inline]
```

Vector of static rules (cells to iterate).

Definition at line 354 of file support-meat.hpp.

7.27.3.14 get_rules_dyn()

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

Rules< Array_Type, Data_Rule_Dyn_Type > * Support< Array_Type, Data_Counter_Type, Data_Rule

_Type, Data_Rule_Dyn_Type >::get_rules_dyn [inline]
```

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

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

7.27.3.15 init_support()

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

160 Class Documentation

7.27.3.16 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.

7.27.3.17 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.

7.27.3.18 reset_array() [2/2]

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

7.27.3.19 set counters()

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

7.27.3.20 set_rules()

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

7.27.3.21 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.

7.27.4 Member Data Documentation

7.27.4.1 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.

7.27.4.2 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.

7.27.4.3 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.

162 Class Documentation

7.27.4.4 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.

7.27.4.5 delete counters

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

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

7.27.4.6 delete_rules

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

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

7.27.4.7 delete_rules_dyn

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

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

7.27.4.8 M

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

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

7.27.4.9 max_num_elements

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

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

7.27.4.10 N

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

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

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

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

7.28 vecHasher < T > Struct Template Reference

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

Public Member Functions

std::size t operator() (std::vector< T > const &dat) const noexcept

7.28.1 Detailed Description

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

Definition at line 86 of file typedefs.hpp.

7.28.2 Member Function Documentation

7.28.2.1 operator()()

Definition at line 87 of file typedefs.hpp.

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

• include/barry/typedefs.hpp

164 Class Documentation

Chapter 8

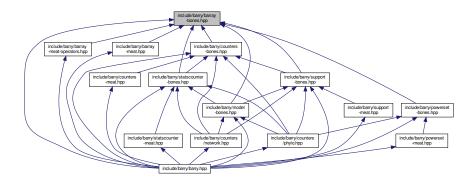
File Documentation

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

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



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



Classes

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

Macros

• #define BARRAY_BONES_HPP 1

8.1.1 Macro Definition Documentation

8.1.1.1 BARRAY_BONES_HPP

#define BARRAY_BONES_HPP 1

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

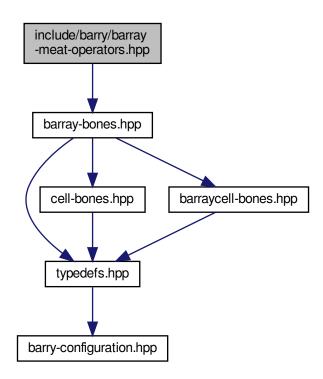
8.2 include/barry/barray-iterator.hpp File Reference

Classes

class ConstBArrayRowIter< Cell_Type, Data_Type >

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

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



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



Macros

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

Functions

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

8.3.1 Macro Definition Documentation

8.3.1.1 BARRY_BARRAY_MEAT_OPERATORS_HPP

```
#define BARRY_BARRAY_MEAT_OPERATORS_HPP 1
```

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

8.3.1.2 COL

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

8.3.1.3 ROW

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

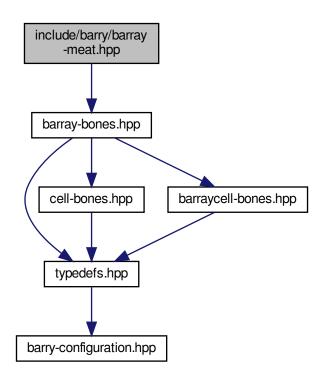
8.3.2 Function Documentation

8.3.2.1 checkdim_()

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

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

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



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



Macros

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

8.4.1 Macro Definition Documentation

8.4.1.1 COL

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

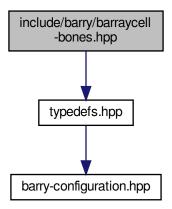
8.4.1.2 ROW

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

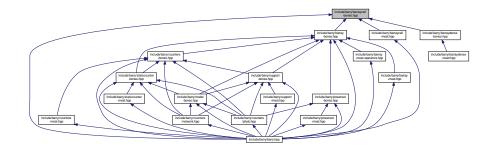
Definition at line 7 of file barray-meat.hpp.

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

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



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

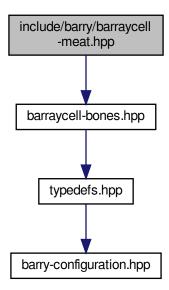


Classes

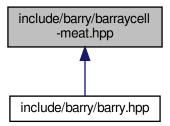
- class BArrayCell
 Cell_Type, Data_Type
- class BArrayCell_const< Cell_Type, Data_Type >

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

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



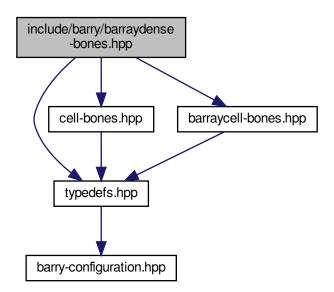
This graph shows which files directly or indirectly include this file:



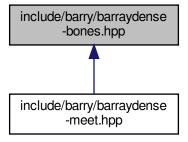
8.7 include/barry/barraydense-bones.hpp File Reference

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

Include dependency graph for barraydense-bones.hpp:



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



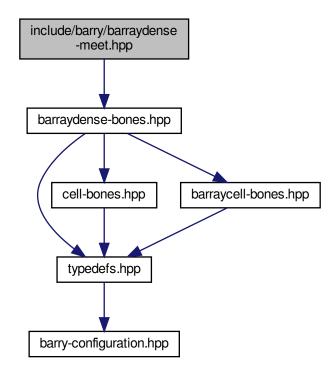
Classes

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

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

#include "barraydense-bones.hpp"

Include dependency graph for barraydense-meet.hpp:



Macros

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

8.8.1 Macro Definition Documentation

8.8.1.1 BARRY_BARRAYDENSE_MEAT_HPP

#define BARRY_BARRAYDENSE_MEAT_HPP

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

8.8.1.2 COL

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

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

8.8.1.3 POS

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

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

8.8.1.4 ROW

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

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

8.8.1.5 ZERO_CELL

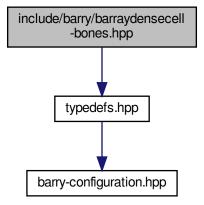
```
#define ZERO_CELL Cell Cell_Type >(static_cast< Cell_Type >(0.0))
```

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

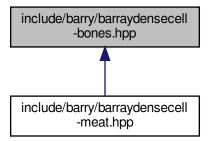
8.9 include/barry/barraydensecell-bones.hpp File Reference

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

Include dependency graph for barraydensecell-bones.hpp:



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

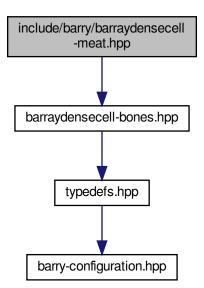


Classes

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

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

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



Macros

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

8.10.1 Macro Definition Documentation

8.10.1.1 BARRY_BARRAYDENSECELL_MEAT_HPP

```
#define BARRY_BARRAYDENSECELL_MEAT_HPP 1
```

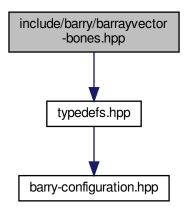
Definition at line 4 of file barraydensecell-meat.hpp.

8.10.1.2 POS

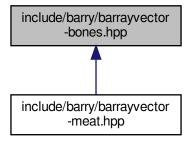
Definition at line 6 of file barraydensecell-meat.hpp.

8.11 include/barry/barrayvector-bones.hpp File Reference

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



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

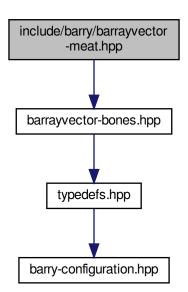


Classes

- class BArrayVector < Cell_Type, Data_Type >
 Row or column of a BArray
- class BArrayVector_const< Cell_Type, Data_Type >

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

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



Macros

#define BARRY_BARRAYVECTOR_MEAT_HPP 1

8.12.1 Macro Definition Documentation

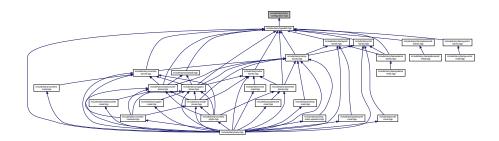
8.12.1.1 BARRY BARRAYVECTOR MEAT HPP

#define BARRY_BARRAYVECTOR_MEAT_HPP 1

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

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

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



Configuration MACROS

These are mostly related to performance. The definitions follow:

- BARRY_USE_UNORDERED_MAP If specified, then barry is compiled using std::unordered_map. Otherwise it will use std::map for the arrays.
- BARRY_USE_SAFE_EXP When specified, it will multiply all likelihoods in Model by (1/-100)/(1/-100) so that numerical overflows are avoided.
- BARRY_USE_ISFINITE When specified, it will introduce a macro that checks whether the likelihood is finite or not.
- printf_barry If not specified, will be defined as printf.
- #define BARRY_SAFE_EXP -100.0
- #define BARRY_ISFINITE(a)
- #define BARRY CHECK SUPPORT(x, maxs)
- #define printf barry printf
- #define BARRY_MAX_NUM_ELEMENTS static_cast< unsigned int >(UINT_MAX/2u)
- template<typename Ta , typename Tb >
 using Map = std::map< Ta, Tb >

8.13.1 Macro Definition Documentation

8.13.1.1 BARRY_CHECK_SUPPORT

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

Definition at line 45 of file barry-configuration.hpp.

8.13.1.2 BARRY_ISFINITE

Definition at line 38 of file barry-configuration.hpp.

8.13.1.3 BARRY_MAX_NUM_ELEMENTS

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

Definition at line 53 of file barry-configuration.hpp.

8.13.1.4 BARRY_SAFE_EXP

```
#define BARRY_SAFE_EXP -100.0
```

Definition at line 31 of file barry-configuration.hpp.

8.13.1.5 printf_barry

```
#define printf_barry printf
```

Definition at line 49 of file barry-configuration.hpp.

8.13.2 Typedef Documentation

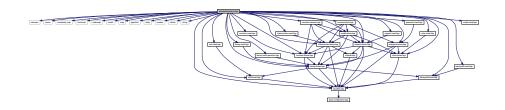
8.13.2.1 Map

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

Definition at line 25 of file barry-configuration.hpp.

8.14 include/barry/barry.hpp File Reference

```
#include <iostream>
#include <vector>
#include <unordered_map>
#include <functional>
#include <stdexcept>
#include <cmath>
#include <map>
#include <algorithm>
#include <utility>
#include <random>
#include <climits>
#include <string>
#include "typedefs.hpp"
#include "cell-bones.hpp"
#include "cell-meat.hpp"
#include "barray-bones.hpp"
#include "barraycell-bones.hpp"
#include "barray-meat.hpp"
#include "barraycell-meat.hpp"
#include "barray-meat-operators.hpp"
#include "counters-bones.hpp"
#include "counters-meat.hpp"
#include "statscounter-bones.hpp"
#include "statscounter-meat.hpp"
#include "support-bones.hpp"
#include "support-meat.hpp"
#include "powerset-bones.hpp"
#include "powerset-meat.hpp"
#include "model-bones.hpp"
#include "model-meat.hpp"
#include "rules-bones.hpp"
#include "rules-meat.hpp"
#include "counters/network.hpp"
#include "counters/phylo.hpp"
Include dependency graph for barry.hpp:
```



Namespaces

barry

barry: Your go-to motif accountant

· barry::counters

Tree class and Treelterator class.

- barry::counters::network
- · barry::counters::phylo

Macros

- #define BARRY_HPP
- #define BARRY_VERSION 0.1
- #define COUNTER FUNCTION(a)
- #define COUNTER_LAMBDA(a)
- #define RULE_FUNCTION(a)
- #define RULE_LAMBDA(a)

8.14.1 Macro Definition Documentation

8.14.1.1 BARRY_HPP

```
#define BARRY_HPP
```

Definition at line 19 of file barry.hpp.

8.14.1.2 BARRY_VERSION

```
#define BARRY_VERSION 0.1
```

Definition at line 21 of file barry.hpp.

8.14.1.3 COUNTER_FUNCTION

```
\begin{tabular}{ll} \# define & COUNTER\_FUNCTION ( \\ & a \end{tabular} \label{eq:counter_function}
```

Value:

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

Definition at line 72 of file barry.hpp.

8.14.1.4 COUNTER_LAMBDA

Definition at line 75 of file barry.hpp.

8.14.1.5 RULE_FUNCTION

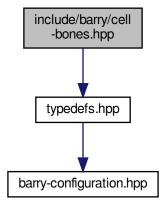
Definition at line 79 of file barry.hpp.

Definition at line 82 of file barry.hpp.

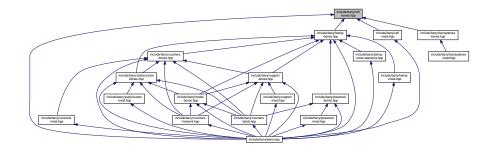
8.14.1.6 RULE_LAMBDA

8.15 include/barry/cell-bones.hpp File Reference

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



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

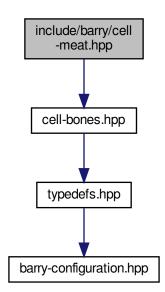


Classes

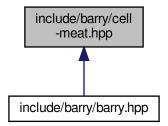
class Cell
 Cell_Type >
 Entries in BArray. For now, it only has two members:

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

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



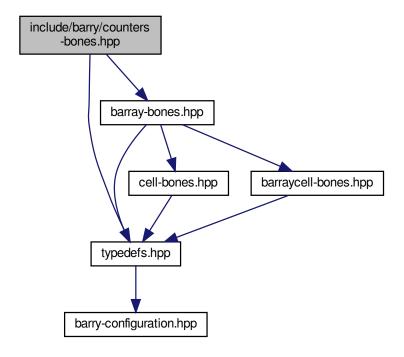
This graph shows which files directly or indirectly include this file:



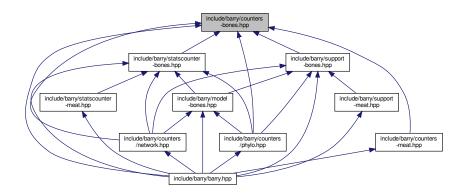
8.17 include/barry/col-bones.hpp File Reference

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

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



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



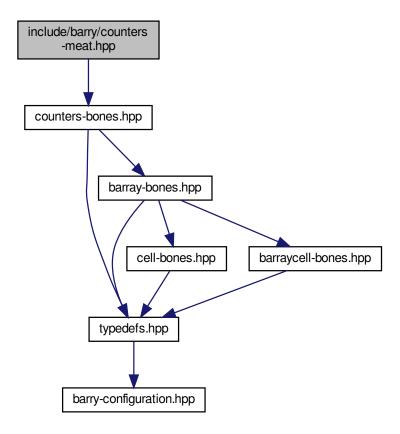
Classes

- class Counter < Array_Type, Data_Type >
 A counter function based on change statistics.

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

#include "counters-bones.hpp"

Include dependency graph for counters-meat.hpp:



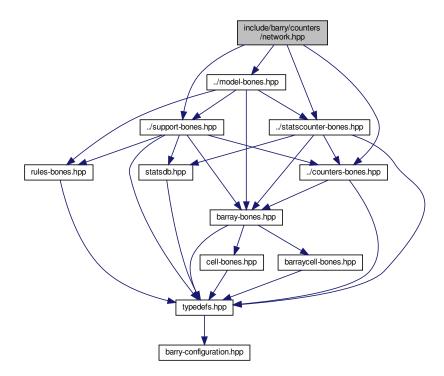
This graph shows which files directly or indirectly include this file:



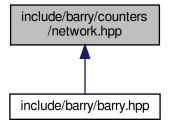
8.20 include/barry/counters/network.hpp File Reference

```
#include "../counters-bones.hpp"
#include "../support-bones.hpp"
```

```
#include "../statscounter-bones.hpp"
#include "../model-bones.hpp"
Include dependency graph for network.hpp:
```



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



Classes

class NetworkData

Data class for Networks.

· class NetCounterData

Data class used to store arbitrary uint or double vectors.

Macros

- #define NET_C_DATA_IDX(i) (data->indices[i])
- #define NET_C_DATA_NUM(i) (data->numbers[i])

Macros for defining counters

- #define NETWORK_COUNTER(a)
- #define NETWORK_COUNTER_LAMBDA(a)

Macros for defining rules

- #define NETWORK_RULE(a)
- #define NETWORK_RULE_LAMBDA(a)

Typedefs

Convenient typedefs for network objects.

- typedef BArray< double, NetworkData > Network
- typedef Counter
 NetWork, NetCounterData > NetCounter
- typedef Counters < Network, NetCounterData > NetCounters
- typedef Support < Network, NetCounterData > NetSupport
- typedef StatsCounter< Network, NetCounterData > NetStatsCounter
- typedef Model < Network, NetCounterData > NetModel
- typedef Rule < Network, bool > NetRule
- typedef Rules < Network, bool > NetRules

Functions

• void counter_edges (NetCounters *counters)

Number of edges.

void counter_isolates (NetCounters *counters)

Number of isolated vertices.

void counter_mutual (NetCounters *counters)

Number of mutual ties.

- void counter_istar2 (NetCounters *counters)
- void counter ostar2 (NetCounters *counters)
- void counter_ttriads (NetCounters *counters)
- void counter_ctriads (NetCounters *counters)
- void counter_density (NetCounters *counters)
- void counter_idegree15 (NetCounters *counters)
- void counter odegree15 (NetCounters *counters)
- void counter_absdiff (NetCounters *counters, uint attr_id, double alpha=1.0)

Sum of absolute attribute difference between ego and alter.

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

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

- NETWORK_COUNTER (init_single_attr)
- void counter_nodeicov (NetCounters *counters, uint attr_id)
- void counter_nodeocov (NetCounters *counters, uint attr_id)

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

Counts number of vertices with a given in-degree.

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

Counts number of vertices with a given out-degree.

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

Counts number of vertices with a given out-degree.

Rules for network models

Parameters

```
rules | A pointer to a NetRules object (Rules < Network, bool > ).
```

void rules_zerodiag (NetRules *rules)
 Number of edges.

8.20.1 Macro Definition Documentation

8.20.1.1 NET C DATA IDX

Definition at line 79 of file network.hpp.

8.20.1.2 NET_C_DATA_NUM

Definition at line 80 of file network.hpp.

8.20.1.3 NETWORK_COUNTER

Function for definition of a network counter function

Definition at line 101 of file network.hpp.

8.20.1.4 NETWORK_COUNTER_LAMBDA

Lambda function for definition of a network counter function

Definition at line 104 of file network.hpp.

8.20.1.5 NETWORK_RULE

Function for definition of a network counter function

Definition at line 113 of file network.hpp.

8.20.1.6 NETWORK_RULE_LAMBDA

Lambda function for definition of a network counter function

Definition at line 116 of file network.hpp.

8.20.2 Typedef Documentation

8.20.2.1 NetCounter

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

Definition at line 88 of file network.hpp.

8.20.2.2 NetCounters

typedef Counters< Network, NetCounterData> NetCounters

Definition at line 89 of file network.hpp.

8.20.2.3 NetModel

typedef Model<Network, NetCounterData> NetModel

Definition at line 92 of file network.hpp.

8.20.2.4 NetRule

typedef Rule<Network,bool> NetRule

Definition at line 93 of file network.hpp.

8.20.2.5 NetRules

typedef Rules<Network,bool> NetRules

Definition at line 94 of file network.hpp.

8.20.2.6 NetStatsCounter

typedef StatsCounter<Network, NetCounterData> NetStatsCounter

Definition at line 91 of file network.hpp.

8.20.2.7 NetSupport

typedef Support<Network, NetCounterData > NetSupport

Definition at line 90 of file network.hpp.

8.20.2.8 Network

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

Definition at line 87 of file network.hpp.

8.20.3 Function Documentation

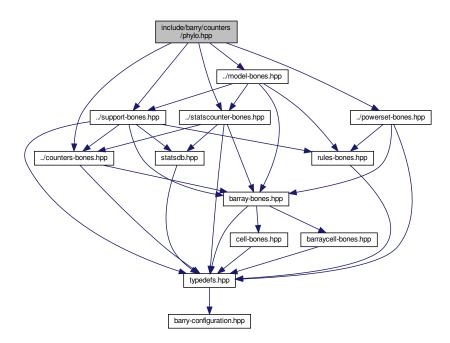
8.20.3.1 rules_zerodiag()

Number of edges.

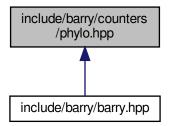
Definition at line 742 of file network.hpp.

8.21 include/barry/counters/phylo.hpp File Reference

```
#include "../counters-bones.hpp"
#include "../support-bones.hpp"
#include "../statscounter-bones.hpp"
#include "../model-bones.hpp"
#include "../powerset-bones.hpp"
Include dependency graph for phylo.hpp:
```



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



Classes

class NodeData

Data definition for the PhyloArray class.

· class PhyloRuleDynData

Macros

- #define PHYLO_COUNTER_LAMBDA(a)
 - Extension of a simple counter.
- #define PHYLO RULE DYN LAMBDA(a)
- #define PHYLO CHECK MISSING()

Typedefs

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

Convenient typedefs for Node objects.

- typedef BArray< uint, NodeData > PhyloArray
- typedef Counter
 PhyloArray, PhyloCounterData > PhyloCounter
- typedef Counters
 PhyloArray, PhyloCounterData > PhyloCounters
- typedef Rule < PhyloArray, PhyloRuleData > PhyloRule
- typedef Rules
 PhyloArray, PhyloRuleData > PhyloRules
- typedef Rule
 PhyloArray, PhyloRuleDynData
 PhyloRuleDyn
- typedef Rules< PhyloArray, PhyloRuleDynData > PhyloRulesDyn
- typedef Support< PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloSupport
- typedef StatsCounter< PhyloArray, PhyloCounterData > PhyloStatsCounter
- typedef Model < PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloModel
- typedef PowerSet< PhyloArray, PhyloRuleData > PhyloPowerSet

Functions

- std::string get last name (bool d)
- void counter_overall_gains (PhyloCounters *counters, bool duplication=true)

Overall functional gains.

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

Functional gains for a specific function (nfun).

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

k genes gain function nfun

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

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

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

Overall functional loss.

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

Cap the number of functions per gene.

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

Total count of losses for an specific function.

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

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

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

Total count of Sub-functionalization events.

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

Co-evolution (joint gain or loss)

void counter longest (PhyloCounters *counters)

Longest branch mutates (either by gain or by loss)

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

Total number of neofunctionalization events.

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

Total number of neofunctionalization events.

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

Function co-opting.

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

Overall functional gains.

8.21.1 Macro Definition Documentation

8.21.1.1 PHYLO_CHECK_MISSING

```
#define PHYLO_CHECK_MISSING( )
```

Value:

```
if (Array.D() == nullptr) \
throw std::logic_error("The array data is nullptr."); \
if (data == nullptr) \
throw std::logic_error("The counter/rule data is nullptr.")
```

Definition at line 94 of file phylo.hpp.

8.21.1.2 PHYLO_COUNTER_LAMBDA

Extension of a simple counter.

It allows specifying extra arguments, in particular, the corresponding sets of rows to which this statistic may be relevant. This could be important in the case of, for example, counting correlation type statistics between function 1 and 2, and between function 1 and 3.

Definition at line 88 of file phylo.hpp.

8.21.1.3 PHYLO_RULE_DYN_LAMBDA

Definition at line 91 of file phylo.hpp.

8.21.2 Typedef Documentation

8.21.2.1 PhyloArray

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

Definition at line 61 of file phylo.hpp.

8.21.2.2 PhyloCounter

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

Definition at line 62 of file phylo.hpp.

8.21.2.3 PhyloCounterData

typedef std::vector< uint > PhyloCounterData

Definition at line 53 of file phylo.hpp.

8.21.2.4 PhyloCounters

typedef Counters< PhyloArray, PhyloCounterData> PhyloCounters

Definition at line 63 of file phylo.hpp.

8.21.2.5 PhyloModel

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

Definition at line 73 of file phylo.hpp.

8.21.2.6 PhyloPowerSet

typedef PowerSet<PhyloArray, PhyloRuleData> PhyloPowerSet

Definition at line 74 of file phylo.hpp.

8.21.2.7 PhyloRule

typedef Rule<PhyloArray,PhyloRuleData> PhyloRule

Definition at line 65 of file phylo.hpp.

8.21.2.8 PhyloRuleData

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

Definition at line 54 of file phylo.hpp.

8.21.2.9 PhyloRuleDyn

```
typedef Rule<PhyloArray,PhyloRuleDynData> PhyloRuleDyn
```

Definition at line 68 of file phylo.hpp.

8.21.2.10 PhyloRules

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

Definition at line 66 of file phylo.hpp.

8.21.2.11 PhyloRulesDyn

```
typedef Rules<PhyloArray,PhyloRuleDynData> PhyloRulesDyn
```

Definition at line 69 of file phylo.hpp.

8.21.2.12 PhyloStatsCounter

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

Definition at line 72 of file phylo.hpp.

8.21.2.13 PhyloSupport

```
typedef Support<PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloSupport
```

Definition at line 71 of file phylo.hpp.

8.21.3 Function Documentation

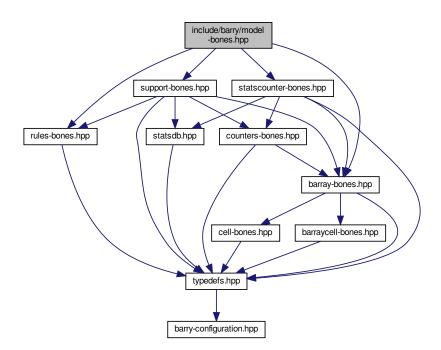
8.21.3.1 get_last_name()

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

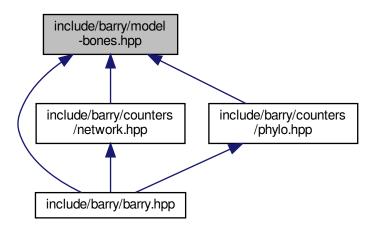
Definition at line 99 of file phylo.hpp.

8.22 include/barry/model-bones.hpp File Reference

```
#include "barray-bones.hpp"
#include "support-bones.hpp"
#include "statscounter-bones.hpp"
#include "rules-bones.hpp"
Include dependency graph for model-bones.hpp:
```



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)

8.22.1 Function Documentation

8.22.1.1 keygen_default()

Array Hasher class (used for computing support)

Definition at line 69 of file model-bones.hpp.

8.22.1.2 likelihood_()

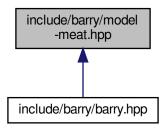
Definition at line 40 of file model-bones.hpp.

8.22.1.3 update_normalizing_constant()

Definition at line 16 of file model-bones.hpp.

8.23 include/barry/model-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



8.24 include/barry/models/geese.hpp File Reference

```
#include "geese/geese-node-bones.hpp"
#include "geese/geese-bones.hpp"
#include "geese/geese-meat.hpp"
#include "geese/geese-meat-constructors.hpp"
#include "geese/geese-meat-likelihood.hpp"
#include "geese/geese-meat-likelihood_exhaust.hpp"
#include "geese/geese-meat-simulate.hpp"
#include "geese/geese-meat-predict.hpp"
#include "geese/flock-bones.hpp"
#include "geese/flock-meet.hpp"
Include dependency graph for geese.hpp:
```



8.25 include/barry/models/geese/flock-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



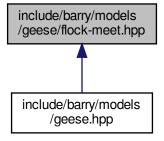
Classes

· class Flock

A Flock is a group of Geese.

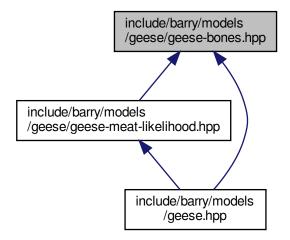
8.26 include/barry/models/geese/flock-meet.hpp File Reference

This graph shows which files directly or indirectly include this file:



8.27 include/barry/models/geese/geese-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



Classes

• class Geese

Annotated Phylo Model.

Macros

• #define INITIALIZED()

Functions

- template<typename Ta , typename Tb > $std::vector < Ta > vector_caster \ (const \ std::vector < Tb > \&x)$
- RULE_FUNCTION (rule_empty_free)
- std::vector< double > keygen_full (const phylocounters::PhyloArray &array)
- bool vec_diff (const std::vector< unsigned int > &s, const std::vector< unsigned int > &a)

8.27.1 Macro Definition Documentation

8.27.1.1 INITIALIZED

```
#define INITIALIZED( )

Value:
    if (!this->initialized) \
        throw std::logic_error("The model has not been initialized yet.");
```

Definition at line 18 of file geese-bones.hpp.

8.27.2 Function Documentation

8.27.2.1 keygen_full()

Definition at line 31 of file geese-bones.hpp.

8.27.2.2 RULE_FUNCTION()

Definition at line 22 of file geese-bones.hpp.

8.27.2.3 vec_diff()

Definition at line 51 of file geese-bones.hpp.

8.27.2.4 vector_caster()

Definition at line 10 of file geese-bones.hpp.

8.28 include/barry/models/geese/geese-meat-constructors.hpp File Reference

This graph shows which files directly or indirectly include this file:



8.29 include/barry/models/geese/geese-meat-likelihood.hpp File Reference

#include "geese-bones.hpp"
Include dependency graph for geese-meat-likelihood.hpp:

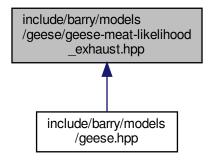


This graph shows which files directly or indirectly include this file:



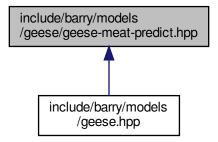
8.30 include/barry/models/geese/geese-meat-likelihood_exhaust.hpp File Reference

This graph shows which files directly or indirectly include this file:



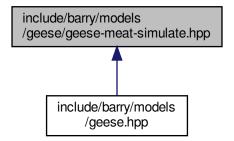
8.31 include/barry/models/geese/geese-meat-predict.hpp File Reference

This graph shows which files directly or indirectly include this file:



8.32 include/barry/models/geese/geese-meat-simulate.hpp File Reference

This graph shows which files directly or indirectly include this file:



8.33 include/barry/models/geese/geese-meat.hpp File Reference

This graph shows which files directly or indirectly include this file:



8.34 include/barry/models/geese/geese-node-bones.hpp File Reference

This graph shows which files directly or indirectly include this file:



Classes

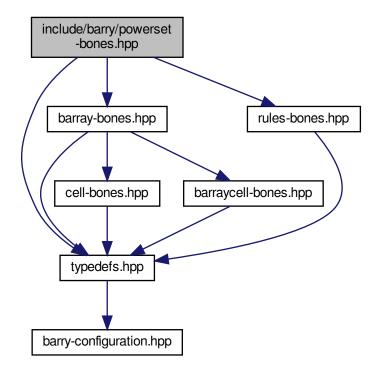
• class Node

A single node for the model.

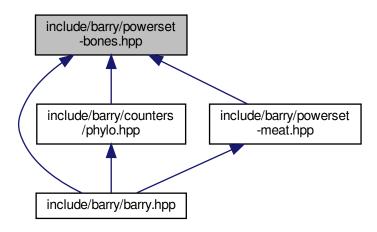
8.35 include/barry/powerset-bones.hpp File Reference

```
#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:

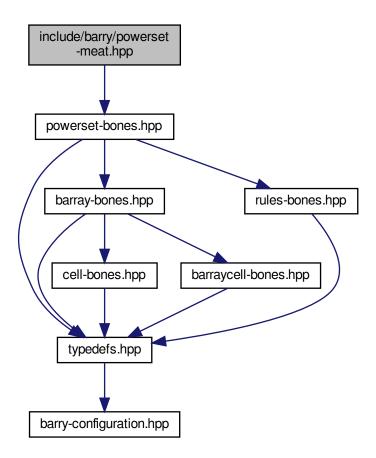


Classes

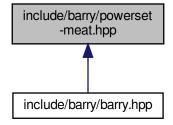
8.36 include/barry/powerset-meat.hpp File Reference

#include "powerset-bones.hpp"

Include dependency graph for powerset-meat.hpp:

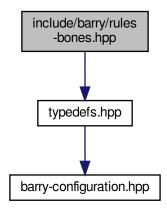


This graph shows which files directly or indirectly include this file:

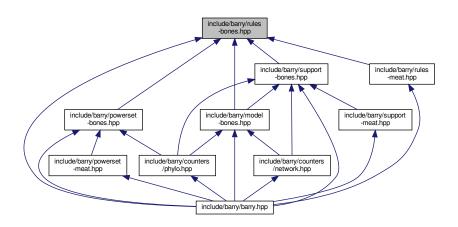


8.37 include/barry/rules-bones.hpp File Reference

#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)

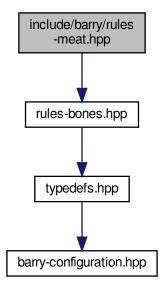
8.37.1 Function Documentation

8.37.1.1 rule_fun_default()

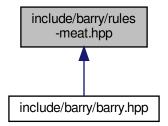
Definition at line 10 of file rules-bones.hpp.

8.38 include/barry/rules-meat.hpp File Reference

```
#include "rules-bones.hpp"
Include dependency graph for rules-meat.hpp:
```



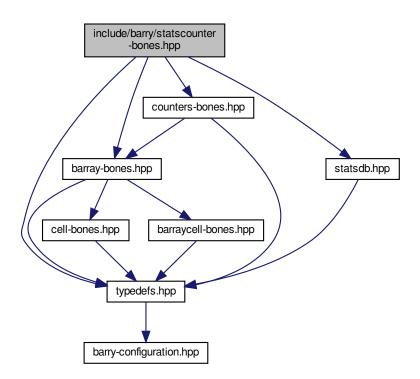
This graph shows which files directly or indirectly include this file:



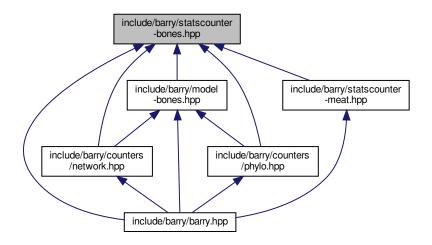
8.39 include/barry/statscounter-bones.hpp File Reference

```
#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:



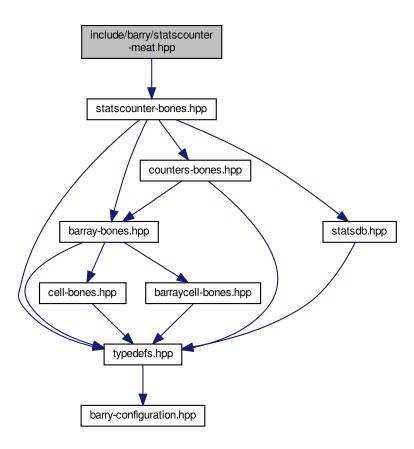
Classes

class StatsCounter< Array_Type, Data_Type >
 Count stats for a single Array.

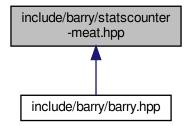
8.40 include/barry/statscounter-meat.hpp File Reference

#include "statscounter-bones.hpp"

Include dependency graph for statscounter-meat.hpp:



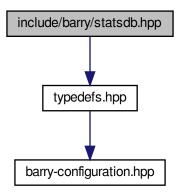
This graph shows which files directly or indirectly include this file:



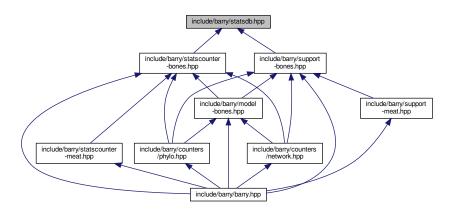
8.41 include/barry/statsdb.hpp File Reference

#include "typedefs.hpp"

Include dependency graph for statsdb.hpp:



This graph shows which files directly or indirectly include this file:



Classes

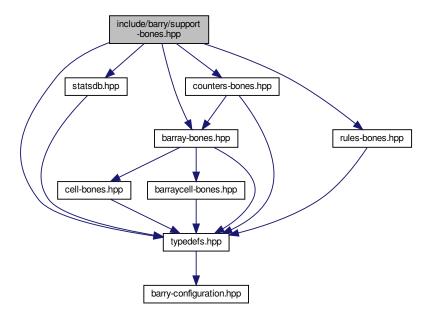
class FreqTable < T >

Database of statistics.

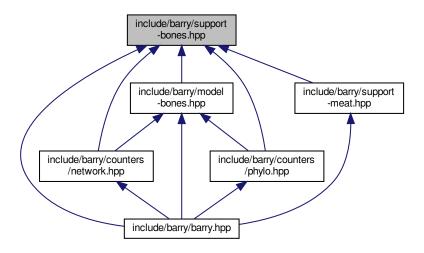
8.42 include/barry/support-bones.hpp File Reference

```
#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:

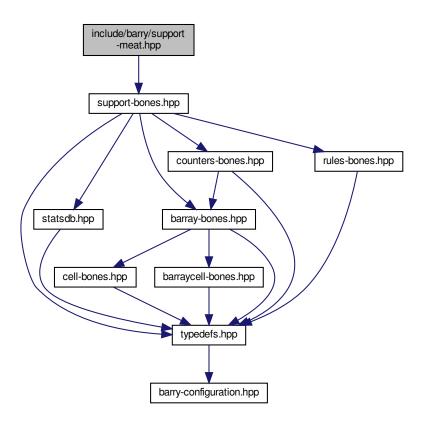


Classes

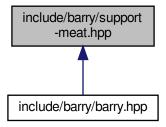
class Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >
 Compute the support of sufficient statistics.

8.43 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

8.43.1 Macro Definition Documentation

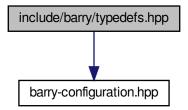
8.43.1.1 BARRY SUPPORT MEAT HPP

#define BARRY_SUPPORT_MEAT_HPP 1

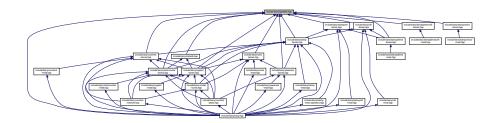
Definition at line 4 of file support-meat.hpp.

8.44 include/barry/typedefs.hpp File Reference

#include "barry-configuration.hpp"
Include dependency graph for typedefs.hpp:



This graph shows which files directly or indirectly include this file:



Classes

- class Entries < Cell_Type >
 - A wrapper class to store source, target, val from a BArray object.
- struct vecHasher< T >

Namespaces

CHECK

Integer constants used to specify which cell should be check.

• EXISTS

Integer constants used to specify which cell should be check to exist or not.

Typedefs

```
• typedef unsigned int uint
```

```
• typedef std::vector< std::pair< std::vector< double >, uint > > Counts_type
```

```
    template<typename Cell_Type >
        using Row_type = Map< uint, Cell< Cell_Type > >
    template<typename Cell_Type >
        using Col_type = Map< uint, Cell< Cell_Type > * >
```

```
• template<typename Ta = double, typename Tb = uint> using MapVec_type = std::unordered_map< std::vector< Ta >, Tb, vecHasher< Ta >>
```

```
    template<typename Array_Type , typename Data_Type >
        using Counter_fun_type = std::function< double(const Array_Type &, uint, uint, Data_Type *)>
        Counter and rule functions.
```

```
    template<typename Array_Type , typename Data_Type >
        using Rule_fun_type = std::function< bool(const Array_Type &, uint, uint, Data_Type *)>
```

Functions

```
   template < typename T > bool vec_equal (const std::vector < T > &a, const std::vector < T > &b)
```

```
Compares if -a- and -b- are equal.

• template<typename T >
bool vec equal approx (const std::vector< T > &a, const std::vector< T > &b, double eps=1e-10)
```

Variables

```
const int CHECK::BOTH = -1
const int CHECK::NONE = 0
const int CHECK::ONE = 1
const int CHECK::TWO = 2
const int EXISTS::BOTH = -1
const int EXISTS::NONE = 0
const int EXISTS::ONE = 1
const int EXISTS::TWO = 1
```

- const int EXISTS::UKNOWN = -1
- const int EXISTS::AS_ZERO = 0
- const int EXISTS::AS_ONE = 1

8.44.1 Typedef Documentation

8.44.1.1 Col type

```
template<typename Cell_Type >
using Col_type = Map< uint, Cell<Cell_Type>* >
```

Definition at line 51 of file typedefs.hpp.

8.44.1.2 Counter_fun_type

```
template<typename Array_Type , typename Data_Type >
using Counter_fun_type = std::function<double(const Array_Type &, uint, uint, Data_Type *)>
```

Counter and rule functions.

Parameters

Array_Type	a BArray
unit,uint	Focal cell
Data_Type	Data associated with the function, for example, id of the attribute in the Array.

Returns

```
Counter_fun_type a double (the change statistic)
Rule_fun_type a bool. True if the cell is blocked.
```

Definition at line 123 of file typedefs.hpp.

8.44.1.3 Counts_type

```
typedef std::vector< std::pair< std::vector<double>, uint > > Counts_type
```

Definition at line 44 of file typedefs.hpp.

8.44.1.4 MapVec_type

```
template<typename Ta = double, typename Tb = uint>
using MapVec_type = std::unordered_map< std::vector< Ta >, Tb, vecHasher<Ta> >
```

Definition at line 105 of file typedefs.hpp.

8.44.1.5 Row_type

```
template<typename Cell_Type >
using Row_type = Map< uint, Cell<Cell_Type> >
```

Definition at line 48 of file typedefs.hpp.

8.44.1.6 Rule_fun_type

```
template<typename Array_Type , typename Data_Type >
using Rule_fun_type = std::function<bool(const Array_Type &, uint, uint, Data_Type *)>
```

Definition at line 126 of file typedefs.hpp.

8.44.1.7 uint

```
typedef unsigned int uint
```

Definition at line 10 of file typedefs.hpp.

8.44.2 Function Documentation

8.44.2.1 vec_equal()

Compares if -a- and -b- are equal.

Parameters

```
a,b Two vectors of the same length
```

Returns

true if all elements are equal.

Definition at line 137 of file typedefs.hpp.

8.44.2.2 vec_equal_approx()

Definition at line 155 of file typedefs.hpp.

8.44.2.3 vec_inner_prod()

Definition at line 175 of file typedefs.hpp.

8.45 README.md File Reference

Index

```
\simBArray
                                                      \simRule
                                                          Rule < Array_Type, Data_Type >, 144
    BArray< Cell Type, Data Type >, 31
\simBArrayCell
                                                      \simRules
    BArrayCell< Cell_Type, Data_Type >, 41
                                                           Rules < Array_Type, Data_Type >, 146
~BArrayCell const
                                                      \simStatsCounter
    BArrayCell_const< Cell_Type, Data_Type >, 44
                                                          StatsCounter< Array_Type, Data_Type >, 150
{\sim}\mathsf{BArrayDense}
                                                      \simSupport
    BArrayDense < Cell_Type, Data_Type >, 50
                                                          Support<
                                                                       Array_Type,
                                                                                      Data_Counter_Type,
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
\simBArrayDenseCell
    BArrayDenseCell< Cell_Type, Data_Type >, 62
~BArrayDenseCell const
                                                      add
    BArrayDenseCell const< Cell Type, Data Type
                                                          Cell< Cell_Type >, 76, 77
         >, 65
                                                          FreqTable < T >, 99
\simBArrayVector
                                                      add array
    BArrayVector< Cell_Type, Data_Type >, 68
                                                          Model<
                                                                      Array_Type,
                                                                                      Data Counter Type,
~BArrayVector const
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
    BArrayVector_const< Cell_Type, Data_Type >, 71
\simCell
                                                      add counter
    Cell< Cell_Type >, 75
                                                          Counters < Array_Type, Data_Type >, 88
\simConstBArrayRowIter
                                                          Model<
                                                                      Array_Type,
                                                                                      Data_Counter_Type,
    ConstBArrayRowlter< Cell Type, Data Type >, 80
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
\simCounter
                                                               115
    Counter< Array_Type, Data_Type >, 83
                                                          StatsCounter< Array Type, Data Type >, 150
\simCounters
                                                          Support<
                                                                       Array Type,
                                                                                      Data Counter Type,
    Counters < Array_Type, Data_Type >, 87
                                                               Data Rule Type, Data Rule Dyn Type >,
\simEntries
    Entries < Cell_Type >, 91
                                                      add data
\simFlock
                                                          Flock, 94
    Flock, 94
                                                      add rule
\simFreqTable
                                                                      Array_Type,
                                                          Model<
                                                                                      Data Counter Type,
    FreqTable < T >, 99
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
\simGeese
                                                               115, 116
    Geese, 104
                                                          PowerSet < Array Type, Data Rule Type >, 139
\simModel
                                                          Rules < Array Type, Data Type >, 147
    Model<
                Array Type,
                                Data Counter Type,
                                                                                      Data_Counter_Type,
                                                          Support<
                                                                       Array_Type,
         Data_Rule_Type, Data_Rule_Dyn_Type >,
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
         114
                                                               156, 157
\simNetCounterData
                                                      add rule dyn
    NetCounterData, 124
                                                          Model<
                                                                      Array_Type,
                                                                                      Data_Counter_Type,
\simNetworkData
                                                               Data_Rule_Type, Data_Rule_Dyn_Type >,
    NetworkData, 126
                                                               116, 117
\simNode
                                                          Support<
                                                                       Array_Type,
                                                                                      Data_Counter_Type,
    Node, 129
                                                               Data Rule Type, Data Rule Dyn Type >,
\simNodeData
                                                               157
    NodeData, 134
                                                      annotations
\simPhyloRuleDynData
                                                          Node, 130
    PhyloRuleDynData, 135
                                                      Arrav
\simPowerSet
                                                          ConstBArrayRowIter< Cell_Type, Data_Type >, 80
    PowerSet < Array_Type, Data_Rule_Type >, 139
                                                      array
```

Node, 130	COL, 168
arrays	ROW, 168
Node, 131	barray-meat.hpp
AS_ONE	COL, 170
EXISTS, 25	ROW, 170
as_vector	BARRAY_BONES_HPP
FreqTable $<$ T $>$, 99	barray-bones.hpp, 166
AS_ZERO	BArrayCell
EXISTS, 25	BArrayCell< Cell Type, Data Type >, 41
	BArrayCell< Cell_Type, Data_Type >, 41
BArray	~BArrayCell, 41
BArray< Cell_Type, Data_Type >, 30, 31	BArray Cell_Type, Data_Type >, 40
BArray< Cell_Type, Data_Type >, 27	BArrayCell, 41
\sim BArray, 31	BArrayDense< Cell_Type, Data_Type >, 60
BArray, 30, 31	operator Cell_Type, 42
BArrayCell< Cell_Type, Data_Type >, 40	operator*=, 42
BArrayCell_const< Cell_Type, Data_Type >, 40	operator+=, 42
clear, 31	operator-=, 42
col, 31	operator/=, 42
D, 32	operator=, 42
default_val, 32	operator==, 43
get_cell, 32	BArrayCell_const
get_col_vec, 32	BArrayCell_const < Cell_Type, Data_Type >, 44
get_entries, 33	
get_row_vec, 33	BArray Cell_const < Cell_Type, Data_Type >, 43
insert_cell, 33, 34	~BArrayCell_const, 44
is_empty, 34	BArray Cell_Type, Data_Type >, 40
ncol, 34	BArrayCell_const, 44
nnozero, 34	BArrayDense < Cell_Type, Data_Type >, 60
nrow, 34	operator Cell_Type, 44
operator*=, 35	operator!=, 44
operator(), 35	operator<, 44
operator+=, 35	operator<=, 45
•	operator>, 45
operator-=, 36	operator>=, 45
operator/=, 36 operator=, 36	operator==, 45
•	BArrayDense
operator==, 37	BArrayDense < Cell_Type, Data_Type >, 48–50
out_of_range, 37	BArrayDense< Cell_Type, Data_Type >, 46
print, 37	\sim BArrayDense, 50
reserve, 37	BArrayCell< Cell_Type, Data_Type >, 60
resize, 37	BArrayCell_const< Cell_Type, Data_Type >, 60
rm_cell, 37	BArrayDense, 48–50
row, 38	clear, 50
set_data, 38	col, 50
swap_cells, 38	D, 51
swap_cols, 38	default_val, 51
swap_rows, 39	get_cell, 51
toggle_cell, 39	get_col_vec, 51, 52
toggle_lock, 39	get_entries, 52
transpose, 39	get_row_vec, 52
visited, 40	insert_cell, 53
zero_col, 39	is_empty, 53
zero_row, 40	ncol, 54
barray-bones.hpp	nnozero, 54
BARRAY_BONES_HPP, 166	nrow, 54
barray-meat-operators.hpp	operator*=, 55
BARRY_BARRAY_MEAT_OPERATORS_HPP,	operator(), 54
168	operator+=, 55
checkdim_, 168	, ,

operator-=, 55, 56	BArrayVector, 67
operator/=, 56	begin, 68
operator=, 56	end, 68
operator==, 57	is_col, 68
out_of_range, 57	is_row, 68
print, 57	operator std::vector< Cell_Type >, 69
reserve, 57	operator*=, 69
resize, 57	operator+=, 69
rm_cell, 58	operator-=, 69
row, 58	operator/=, 69
set_data, 58	operator=, 70
swap_cells, 59	operator==, 70
swap_cols, 59	size, 70
swap_rows, 59	barrayvector-meat.hpp
toggle_cell, 59	BARRY_BARRAYVECTOR_MEAT_HPP, 179
toggle_lock, 59	BArrayVector_const
transpose, 60	BArrayVector_const< Cell_Type, Data_Type >, 71
visited, 61	BArrayVector_const< Cell_Type, Data_Type >, 70
zero col, 60	~BArrayVector_const, 71
zero_row, 60	BArrayVector_const, 71
barraydense-meet.hpp	begin, 72
BARRY_BARRAYDENSE_MEAT_HPP, 174	end, 72
COL, 174	is_col, 72
POS, 175	is_row, 72
ROW, 175	operator std::vector< Cell_Type >, 72
ZERO_CELL, 175	operator!=, 72
BArrayDenseCell	operator<, 73
BArrayDenseCell< Cell_Type, Data_Type >, 62	operator<=, 73
BArrayDenseCell< Cell_Type, Data_Type >, 61	operator>, 73
~BArrayDenseCell, 62	operator>=, 73
BArrayDenseCell, 62	operator==, 73
operator Cell_Type, 62	size, 74
operator*=, 62	barry, 23
operator+=, 63	barry-configuration.hpp
operator-=, 63	BARRY_CHECK_SUPPORT, 180
operator/=, 63	BARRY_ISFINITE, 180
•	BARRY_MAX_NUM_ELEMENTS, 180
operator=, 63 operator==, 63	BARRY_SAFE_EXP, 180
barraydensecell-meat.hpp	Map, 180
• • • • • • • • • • • • • • • • • • • •	•
BARRY_BARRAYDENSECELL_MEAT_HPP, 177	printf_barry, 180 barry.hpp
POS, 177 BArrayDenseCell const	BARRY_HPP, 182
BArrayDenseCell const< Cell Type, Data Type	
	BARRY_VERSION, 182 COUNTER FUNCTION, 182
>, 64	-
BArrayDenseCell_const< Cell_Type, Data_Type >, 64	COUNTER_LAMBDA, 182
~BArrayDenseCell_const, 65	RULE_FUNCTION, 183
BArrayDenseCell_const, 64	RULE_LAMBDA, 183
operator Cell_Type, 65	barry::counters, 23
operator!=, 65	barry::counters::network, 24
operator<, 65	barry::counters::phylo, 24
operator<=, 65	BARRY_BARRAY_MEAT_OPERATORS_HPP
operator>, 66	barray-meat-operators.hpp, 168
operator>=, 66	BARRY_BARRAYDENSE_MEAT_HPP
operator==, 66	barraydense-meet.hpp, 174
BArrayVector	BARRY_BARRAYDENSECELL_MEAT_HPP
BArrayVector< Cell_Type, Data_Type >, 67	barraydensecell-meat.hpp, 177
BArrayVector< Cell_Type, Data_Type >, 66	BARRY_BARRAYVECTOR_MEAT_HPP
\sim BArrayVector, 68	barrayvector-meat.hpp, 179

BARRY_CHECK_SUPPORT	Counters < Array_Type, Data_Type >, 88
barry-configuration.hpp, 180	FreqTable < T >, 100
BARRY_HPP	Rules < Array_Type, Data_Type >, 147
barry.hpp, 182	COL
BARRY_ISFINITE	barray-meat-operators.hpp, 168
barry-configuration.hpp, 180	barray-meat.hpp, 170
BARRY_MAX_NUM_ELEMENTS	barraydense-meet.hpp, 174
barry-configuration.hpp, 180	col
BARRY_SAFE_EXP	BArray< Cell_Type, Data_Type >, 31
barry-configuration.hpp, 180	BArrayDense< Cell_Type, Data_Type >, 50
BARRY_SUPPORT_MEAT_HPP	Col_type
support-meat.hpp, 220	typedefs.hpp, 222
BARRY_VERSION	ConstBArrayRowlter
barry.hpp, 182	ConstBArrayRowlter< Cell_Type, Data_Type >, 79
begin	ConstBArrayRowlter< Cell_Type, Data_Type >, 79
BArrayVector< Cell_Type, Data_Type >, 68	∼ConstBArrayRowIter, 80
BArrayVector_const< Cell_Type, Data_Type >, 72	Array, 80
PowerSet< Array_Type, Data_Rule_Type >, 139	ConstBArrayRowlter, 79
blengths	current_col, 80
NodeData, 134	current_row, 80
BOTH	iter, 80
CHECK, 24	coordinates_free
EXISTS, 25	PowerSet < Array_Type, Data_Rule_Type >, 141
calc	Support< Array_Type, Data_Counter_Type
PowerSet< Array_Type, Data_Rule_Type >, 140	Data_Rule_Type, Data_Rule_Dyn_Type >
Support< Array_Type, Data_Counter_Type,	161
Data_Rule_Type, Data_Rule_Dyn_Type >,	coordinates_locked
157	PowerSet< Array_Type, Data_Rule_Type >, 141
calc_reduced_sequence	Support< Array_Type, Data_Counter_Type
Geese, 104	Data_Rule_Type, Data_Rule_Dyn_Type >
calc_sequence	161
Geese, 104	Country Array Type Data Type > 92
Cell	Counter< Array_Type, Data_Type >, 83
Cell< Cell_Type >, 75, 76	count_all StatsCounter< Array_Type, Data_Type >, 151
Cell< Cell_Type >, 74	count_current
~Cell, 75	Count Current
	StatsCounter< Array_Type, Data_Type >, 151
add, 76, 77 Cell, 75, 76	StatsCounter< Array_Type, Data_Type >, 151 count_fun
add, 76, 77 Cell, 75, 76	StatsCounter< Array_Type, Data_Type >, 151 count_fun Counter< Array_Type, Data_Type >, 84
add, 76, 77	StatsCounter< Array_Type, Data_Type >, 151 count_fun Counter< Array_Type, Data_Type >, 84 count_init
add, 76, 77 Cell, 75, 76 operator Cell_Type, 77	StatsCounter< Array_Type, Data_Type >, 151 count_fun Counter< Array_Type, Data_Type >, 84 count_init StatsCounter< Array_Type, Data_Type >, 151
add, 76, 77 Cell, 75, 76 operator Cell_Type, 77 operator!=, 77	StatsCounter< Array_Type, Data_Type >, 151 count_fun Counter< Array_Type, Data_Type >, 84 count_init StatsCounter< Array_Type, Data_Type >, 151 Counter
add, 76, 77 Cell, 75, 76 operator Cell_Type, 77 operator!=, 77 operator=, 77, 78	StatsCounter< Array_Type, Data_Type >, 151 count_fun Counter< Array_Type, Data_Type >, 84 count_init StatsCounter< Array_Type, Data_Type >, 151 Counter Counter< Array_Type, Data_Type >, 82, 83
add, 76, 77 Cell, 75, 76 operator Cell_Type, 77 operator!=, 77 operator==, 77, 78 operator==, 78	StatsCounter< Array_Type, Data_Type >, 151 count_fun Counter< Array_Type, Data_Type >, 84 count_init StatsCounter< Array_Type, Data_Type >, 151 Counter Counter< Array_Type, Data_Type >, 82, 83 Counter< Array_Type, Data_Type >, 81
add, 76, 77 Cell, 75, 76 operator Cell_Type, 77 operator!=, 77 operator=, 77, 78 operator==, 78 value, 78	StatsCounter< Array_Type, Data_Type >, 151 count_fun Counter< Array_Type, Data_Type >, 84 count_init StatsCounter< Array_Type, Data_Type >, 151 Counter Counter< Array_Type, Data_Type >, 82, 83 Counter< Array_Type, Data_Type >, 81 ~Counter, 83
add, 76, 77 Cell, 75, 76 operator Cell_Type, 77 operator!=, 77 operator=, 77, 78 operator==, 78 value, 78 visited, 78	StatsCounter< Array_Type, Data_Type >, 151 count_fun Counter< Array_Type, Data_Type >, 84 count_init StatsCounter< Array_Type, Data_Type >, 151 Counter Counter< Array_Type, Data_Type >, 82, 83 Counter< Array_Type, Data_Type >, 81 ~Counter, 83 count, 83
add, 76, 77 Cell, 75, 76 operator Cell_Type, 77 operator!=, 77 operator=, 77, 78 operator==, 78 value, 78 visited, 78 change_stats	StatsCounter< Array_Type, Data_Type >, 151 count_fun Counter< Array_Type, Data_Type >, 84 count_init StatsCounter< Array_Type, Data_Type >, 151 Counter Counter< Array_Type, Data_Type >, 82, 83 Counter< Array_Type, Data_Type >, 81 ~Counter, 83 count, 83 count_fun, 84
add, 76, 77 Cell, 75, 76 operator Cell_Type, 77 operator!=, 77 operator==, 78, operator==, 78 value, 78 visited, 78 change_stats Support< Array_Type, Data_Counter_Type,	StatsCounter< Array_Type, Data_Type >, 151 count_fun
add, 76, 77 Cell, 75, 76 operator Cell_Type, 77 operator!=, 77 operator==, 77, 78 operator==, 78 value, 78 visited, 78 change_stats Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,	StatsCounter< Array_Type, Data_Type >, 151 count_fun Counter< Array_Type, Data_Type >, 84 count_init StatsCounter< Array_Type, Data_Type >, 151 Counter Counter< Array_Type, Data_Type >, 82, 83 Counter< Array_Type, Data_Type >, 81 ~Counter, 83 count, 83 count_fun, 84 Counter, 82, 83 data, 84
add, 76, 77 Cell, 75, 76 operator Cell_Type, 77 operator!=, 77 operator==, 78 operator==, 78 value, 78 visited, 78 change_stats Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 161	StatsCounter< Array_Type, Data_Type >, 151 count_fun Counter< Array_Type, Data_Type >, 84 count_init StatsCounter< Array_Type, Data_Type >, 151 Counter Counter< Array_Type, Data_Type >, 82, 83 Counter< Array_Type, Data_Type >, 81 ~Counter, 83 count, 83 count_fun, 84 Counter, 82, 83 data, 84 delete_data, 85
add, 76, 77 Cell, 75, 76 operator Cell_Type, 77 operator!=, 77 operator==, 78 operator==, 78 value, 78 visited, 78 change_stats Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 161 CHECK, 24	StatsCounter< Array_Type, Data_Type >, 151 count_fun Counter< Array_Type, Data_Type >, 84 count_init StatsCounter< Array_Type, Data_Type >, 151 Counter Counter< Array_Type, Data_Type >, 82, 83 Counter< Array_Type, Data_Type >, 81 ~Counter, 83 count, 83 count_fun, 84 Counter, 82, 83 data, 84 delete_data, 85 desc, 85
add, 76, 77 Cell, 75, 76 operator Cell_Type, 77 operator!=, 77 operator==, 78 operator==, 78 value, 78 visited, 78 change_stats Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 161 CHECK, 24 BOTH, 24 NONE, 24 ONE, 24	StatsCounter< Array_Type, Data_Type >, 151 count_fun Counter< Array_Type, Data_Type >, 84 count_init StatsCounter< Array_Type, Data_Type >, 151 Counter Counter< Array_Type, Data_Type >, 82, 83 Counter< Array_Type, Data_Type >, 81 ~Counter, 83 count, 83 count_fun, 84 Counter, 82, 83 data, 84 delete_data, 85 desc, 85 init, 83
add, 76, 77 Cell, 75, 76 operator Cell_Type, 77 operator!=, 77 operator==, 78 operator==, 78 value, 78 visited, 78 change_stats Support< Array_Type, Data_Counter_Type,	StatsCounter< Array_Type, Data_Type >, 151 count_fun Counter< Array_Type, Data_Type >, 84 count_init StatsCounter< Array_Type, Data_Type >, 151 Counter Counter< Array_Type, Data_Type >, 82, 83 Counter< Array_Type, Data_Type >, 81 ~Counter, 83 count, 83 count_fun, 84 Counter, 82, 83 data, 84 delete_data, 85 desc, 85 init, 83 init_fun, 85
add, 76, 77 Cell, 75, 76 operator Cell_Type, 77 operator!=, 77 operator==, 78 operator==, 78 value, 78 visited, 78 change_stats Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 161 CHECK, 24 BOTH, 24 NONE, 24 ONE, 24	StatsCounter< Array_Type, Data_Type >, 151 count_fun Counter< Array_Type, Data_Type >, 84 count_init StatsCounter< Array_Type, Data_Type >, 151 Counter Counter< Array_Type, Data_Type >, 82, 83 Counter< Array_Type, Data_Type >, 81 ~Counter, 83 count, 83 count_fun, 84 Counter, 82, 83 data, 84 delete_data, 85 desc, 85 init, 83 init_fun, 85 name, 85
add, 76, 77 Cell, 75, 76 operator Cell_Type, 77 operator!=, 77 operator==, 78 operator==, 78 value, 78 visited, 78 change_stats Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 161 CHECK, 24 BOTH, 24 NONE, 24 ONE, 24 TWO, 24	StatsCounter< Array_Type, Data_Type >, 151 count_fun
add, 76, 77 Cell, 75, 76 operator Cell_Type, 77 operator!=, 77 operator==, 78 operator==, 78 value, 78 visited, 78 change_stats Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 161 CHECK, 24 BOTH, 24 NONE, 24 ONE, 24 TWO, 24 checkdim_	StatsCounter< Array_Type, Data_Type >, 151 count_fun Counter< Array_Type, Data_Type >, 84 count_init StatsCounter< Array_Type, Data_Type >, 151 Counter Counter< Array_Type, Data_Type >, 82, 83 Counter< Array_Type, Data_Type >, 82, 83 Counter< Array_Type, Data_Type >, 81 ~Counter, 83 count, 83 count_fun, 84 Counter, 82, 83 data, 84 delete_data, 85 desc, 85 init, 83 init_fun, 85 name, 85 operator=, 84 counter_absdiff
add, 76, 77 Cell, 75, 76 operator Cell_Type, 77 operator!=, 77 operator==, 78 operator==, 78 value, 78 visited, 78 change_stats Support< Array_Type, Data_Counter_Type,	StatsCounter< Array_Type, Data_Type >, 151 count_fun

Phylo counters, 16	Network counters, 14
counter_cogain	counter_overall_changes
Phylo counters, 16	Phylo counters, 18
counter_ctriads	counter_overall_gains
Network counters, 11	Phylo counters, 19
counter_degree	counter_overall_loss
Network counters, 11	Phylo counters, 19
counter_density	counter_subfun
Network counters, 11	Phylo counters, 19
counter_diff	counter_ttriads
Network counters, 11	Network counters, 14
counter_edges	Counters
Network counters, 12	Counters< Array_Type, Data_Type >, 86, 87
Counter_fun_type	Counters< Array_Type, Data_Type >, 86
typedefs.hpp, 222	∼Counters, 87
COUNTER_FUNCTION	add_counter, 88
barry.hpp, 182	clear, 88
counter_gains	Counters, 86, 87
Phylo counters, 16	operator=, 88, 89
counter_gains_k_offspring	operator[], 89
Phylo counters, 17	size, 90
counter_genes_changing	Counting, 9
Phylo counters, 17	counts
counter_idegree	PhyloRuleDynData, 136
Network counters, 12	Counts_type
counter_idegree15	typedefs.hpp, 222
Network counters, 12	current_col
counter_isolates	ConstBArrayRowlter< Cell_Type, Data_Type >, 80
Network counters, 12	current row
counter_istar2	ConstBArrayRowlter< Cell_Type, Data_Type >, 80
Network counters, 12	current_stats
COUNTER_LAMBDA	Support< Array_Type, Data_Counter_Type,
barry.hpp, 182	Data_Rule_Type, Data_Rule_Dyn_Type >,
counter_longest	161
Phylo counters, 17	
counter_loss	D
Phylo counters, 17	BArray< Cell_Type, Data_Type >, 32
counter_maxfuns	BArrayDense< Cell_Type, Data_Type >, 51
Phylo counters, 18	Rule < Array_Type, Data_Type >, 145
counter_mutual	dat
Network counters, 13	Flock, 97
counter_neofun	data
Phylo counters, 18	Counter< Array_Type, Data_Type >, 84
counter_neofun_a2b	PowerSet < Array_Type, Data_Rule_Type >, 142
Phylo counters, 18	default_val
counter_nodecov	BArray< Cell_Type, Data_Type >, 32
Network counters, 13	BArrayDense < Cell_Type, Data_Type >, 51
counter_nodeicov	delete_counters
Network counters, 13	Support< Array_Type, Data_Counter_Type,
counter_nodematch	Data_Rule_Type, Data_Rule_Dyn_Type >,
Network counters, 13	162
counter_nodeocov	delete_data
Network counters, 13	Counter< Array_Type, Data_Type >, 85
counter_odegree	delete_rengine
Network counters, 14	Geese, 109
counter_odegree15	delete_rules
Network counters, 14	Support< Array_Type, Data_Counter_Type,
counter_ostar2	Data_Rule_Type, Data_Rule_Dyn_Type >,
	162

delete_rules_dyn Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 162 delete_support Geese, 110 desc Counter< Array_Type, Data_Type >, 85 directed NetworkData, 127 duplication Node, 131 NodeData, 134 PhyloRuleDynData, 136	support_size, 97 FreqTable FreqTable < T >, 99 FreqTable < T >, 98
EmptyArray PowerSet < Array_Type, Data_Rule_Type >, 142 end BArrayVector < Cell_Type, Data_Type >, 68 BArrayVector_const < Cell_Type, Data_Type >, 72 PowerSet < Array_Type, Data_Rule_Type >, 140 Entries Entries < Cell_Type >, 91 Entries < Cell_Type >, 90 ~Entries, 91 Entries, 91 resize, 91 source, 92 target, 92 val, 92 EXISTS, 25 AS_ONE, 25 AS_ZERO, 25 BOTH, 25 NONE, 26 ONE, 26 TWO, 26 UKNOWN, 26 Flock, 92 ~Flock, 94 add_data, 94	Geese, 101 ~Geese, 104 calc_reduced_sequence, 104 calc_sequence, 104 delete_rengine, 109 delete_support, 110 Geese, 103, 104 get_counters, 104 get_model, 105 get_probabilities, 105 get_states, 105 get_states, 105 get_support, 105 inherit_support, 105 init, 106 init_node, 106 initialized, 110 likelihood, 106 likelihood_exhaust, 106 map_to_nodes, 110 nfunctions, 110 nfuns, 106 nleafs, 107 nnodes, 107 nodes, 110 nterms, 107 observed_counts, 107
dat, 97 Flock, 93 get_counters, 94 get_support, 94 init, 95 initialized, 97 likelihood_joint, 95 model, 97 nfunctions, 98 nfuns, 95 nleafs, 95 nnodes, 96 nterms, 96 ntrees, 96 operator(), 96 rengine, 98 set_seed, 97	operator=, 107, 108 parse_polytomies, 108 predict, 108 predict_backend, 108 print_observed_counts, 108 reduced_sequence, 110 sequence, 111 set_seed, 109 simulate, 109 support_size, 109 update_annotations, 109 geese-bones.hpp INITIALIZED, 203 keygen_full, 204 RULE_FUNCTION, 204 vec_diff, 204 vector_caster, 204

get_cell	Model < Array_Type, Data_Counter_Type,
BArray< Cell_Type, Data_Type >, 32 BArrayDense< Cell Type, Data Type >, 51	Data_Rule_Type, Data_Rule_Dyn_Type >, 118
get_col_vec	get row vec
BArray< Cell_Type, Data_Type >, 32	BArray< Cell_Type, Data_Type >, 33
BArrayDense < Cell_Type, Data_Type >, 51, 52	BArrayDense< Cell_Type, Data_Type >, 52
get_counters	get_rules
Flock, 94	Model< Array_Type, Data_Counter_Type,
Geese, 104	Data_Rule_Type, Data_Rule_Dyn_Type >,
Model< Array_Type, Data_Counter_Type,	118
Data_Rule_Type, Data_Rule_Dyn_Type >, 117	Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,
StatsCounter< Array_Type, Data_Type >, 151	159
Support< Array_Type, Data_Counter_Type,	
Data_Rule_Type, Data_Rule_Dyn_Type >,	Model < Array_Type, Data_Counter_Type,
158	Data_Rule_Type, Data_Rule_Dyn_Type >,
get_counts	118
Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 158	Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 159
get_counts_ptr	get_seq
Support< Array_Type, Data_Counter_Type,	Rules < Array_Type, Data_Type >, 147
Data_Rule_Type, Data_Rule_Dyn_Type >,	
158	Geese, 105
get_current_stats	get_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 >,
158	118
get_data	get_support
FreqTable < T >, 100	Flock, 94
PowerSet< Array_Type, Data_Rule_Type >, 140 Support< Array_Type, Data_Counter_Type,	Geese, 105 Model< Array_Type, Data_Counter_Type,
Data_Rule_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,	Data_Rule_Type, Data_Rule_Dyn_Type >,
158	118
get_data_ptr	
FreqTable< T >, 100	id
PowerSet < Array_Type, Data_Rule_Type >, 140	Node, 131
get_entries	include/barry/barray-bones.hpp, 165
BArray< Cell_Type, Data_Type >, 33	include/barry/barray-iterator.hpp, 166
BArrayDense < Cell_Type, Data_Type >, 52	include/barry/barray-meat-operators.hpp, 167 include/barry/barray-meat.hpp, 169
get_last_name	include/barry/barraycell-bones.hpp, 170
phylo.hpp, 198	include/barry/barraycell-meat.hpp, 171
get_model Geese, 105	include/barry/barraydense-bones.hpp, 172
get_norm_const	include/barry/barraydense-meet.hpp, 173
Model Array_Type, Data_Counter_Type,	include/barry/barraydensecell-bones.hpp, 175
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/barraydensecell-meat.hpp, 176
117	include/barry/barrayvector-bones.hpp, 177
get_parent	include/barry/barrayvector-meat.hpp, 178
Node, 130	include/barry/barry-configuration.hpp, 179
get_probabilities	include/barry/barry.hpp, 181
Geese, 105	include/barry/cell-bones.hpp, 183
get_pset	include/barry/cell-meat.hpp, 184
Model< Array_Type, Data_Counter_Type,	include/barry/col-bones.hpp, 185
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/counters-bones.hpp, 185
117	include/barry/counters-meat.hpp, 186
get_rengine	include/barry/counters/network.hpp, 187 include/barry/counters/phylo.hpp, 193
Geese, 105	include/barry/model-bones.hpp, 199

include/barry/model-meat.hpp, 201 include/barry/models/geese.hpp, 201 include/barry/models/geese/flock-bones.hpp, 202 include/barry/models/geese/flock-meet.hpp, 202 include/barry/models/geese/geese-bones.hpp, 203 include/barry/models/geese/geese-meat-constructors.hpp 205 include/barry/models/geese/geese-meat-likelihood.hpp, 205	iter ConstBArrayRowIter< Cell_Type, Data_Type >, 80
include/barry/models/geese/geese-meat-likelihood_exhau	g (朝)侯(, detault model-bones.hpp, 200
206 include/barry/models/geese/geese-meat-predict.hpp, 207	keygen_full geese-bones.hpp, 204
include/barry/models/geese/geese-meat-simulate.hpp, 207	lb
include/barry/models/geese/geese-meat.hpp, 208 include/barry/models/geese/geese-node-bones.hpp, 208 include/barry/powerset-bones.hpp, 209 include/barry/powerset-meat.hpp, 210 include/barry/rules-bones.hpp, 212	PhyloRuleDynData, 136 likelihood Geese, 106 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 119
include/barry/rules-meat.hpp, 213	likelihood_ model-bones.hpp, 200
include/barry/statscounter-bones.hpp, 214 include/barry/statscounter-meat.hpp, 215	likelihood_exhaust
include/barry/statsdb.hpp, 216	Geese, 106
include/barry/support-bones.hpp, 217	likelihood_joint Flock, 95
include/barry/support-meat.hpp, 219	likelihood total
include/barry/typedefs.hpp, 220	Model < Array_Type, Data_Counter_Type,
indices NotCounterDate 124	Data_Rule_Type, Data_Rule_Dyn_Type >,
NetCounterData, 124 inherit_support	119
Geese, 105	
init	M
Counter< Array_Type, Data_Type >, 83 Flock, 95 Geese, 106	PowerSet < Array_Type, Data_Rule_Type >, 142 Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 162
init_fun	Map
Counter< Array_Type, Data_Type >, 85	barry-configuration.hpp, 180
init_node Geese, 106	map_to_nodes
init_support	Geese, 110
PowerSet< Array_Type, Data_Rule_Type >, 140	MapVec_type
Support< Array_Type, Data_Counter_Type,	typedefs.hpp, 222 max_num_elements
Data_Rule_Type, Data_Rule_Dyn_Type >,	Support< Array_Type, Data_Counter_Type,
159	Data_Rule_Type, Data_Rule_Dyn_Type >,
INITIALIZED	162
geese-bones.hpp, 203	Model
initialized Flock, 97	Model < Array_Type, Data_Counter_Type,
Geese, 110	Data_Rule_Type, Data_Rule_Dyn_Type >,
insert_cell	113, 114
BArray< Cell_Type, Data_Type >, 33, 34	model
BArrayDense < Cell_Type, Data_Type >, 53	Flock, 97
is_col	Model < Array_Type, Data_Counter_Type, Data_Rule_Type
BArrayVector< Cell_Type, Data_Type >, 68	Data_Rule_Dyn_Type >, 111 ~Model, 114
BArrayVector_const< Cell_Type, Data_Type >, 72	add_array, 114
is_empty	add_counter, 115
BArray< Cell_Type, Data_Type >, 34	add_rule, 115, 116

add_rule_dyn, 116, 117	network.hpp, 192
get_counters, 117	NetRules
get_norm_const, 117	network.hpp, 192
get_pset, 117	NetStatsCounter
get_rengine, 118	network.hpp, 192
get_rules, 118	NetSupport
get_rules_dyn, 118	network.hpp, 192
get_stats, 118	Network
get_support, 118	network.hpp, 192
likelihood, 119	Network counters, 10
likelihood_total, 119	counter_absdiff, 11
Model, 113, 114	counter_ctriads, 11
nterms, 120	counter_degree, 11
operator=, 120	counter_density, 11
print_stats, 120	counter_diff, 11
sample, 120, 121	counter_edges, 12
set_counters, 121	counter_idegree, 12
set_keygen, 121	counter_idegree15, 12
set_rengine, 121	counter_isolates, 12
set_rules, 122	counter_istar2, 12
set_rules_dyn, 122	counter_mutual, 13
set_seed, 122	counter_nodecov, 13
size, 122	counter_nodeicov, 13
size_unique, 122	counter_nodematch, 13
store_psets, 123	counter_nodeocov, 13
support_size, 123	counter_odegree, 14
model-bones.hpp	counter_odegree15, 14
keygen_default, 200	counter_ostar2, 14
likelihood_, 200	counter_ttriads, 14
update_normalizing_constant, 200	NETWORK_COUNTER, 14
	network.hpp
N	NET_C_DATA_IDX, 190
PowerSet < Array_Type, Data_Rule_Type >, 142	NET_C_DATA_NUM, 190
Support< Array_Type, Data_Counter_Type,	NetCounter, 191
Data_Rule_Type, Data_Rule_Dyn_Type >,	NetCounters, 191
163	NetModel, 192
name	NetRule, 192
Counter< Array_Type, Data_Type >, 85	NetRules, 192
narray	NetStatsCounter, 192
Node, 131	NetSupport, 192
ncol	Network, 192
BArray< Cell_Type, Data_Type >, 34	NETWORK_COUNTER, 190
BArrayDense< Cell_Type, Data_Type >, 54	NETWORK_COUNTER_LAMBDA, 190
NET_C_DATA_IDX	NETWORK_RULE, 191
network.hpp, 190	NETWORK_RULE_LAMBDA, 191
NET_C_DATA_NUM	rules zerodiag, 193
network.hpp, 190	NETWORK COUNTER
NetCounter	Network counters, 14
network.hpp, 191	network.hpp, 190
NetCounterData, 123	NETWORK_COUNTER_LAMBDA
\sim NetCounterData, 124	network.hpp, 190
indices, 124	NETWORK_RULE
NetCounterData, 124	network.hpp, 191
numbers, 125	NETWORK_RULE_LAMBDA
NetCounters	network.hpp, 191
network.hpp, 191	NetworkData, 125
NetModel	\sim NetworkData, 126
network.hpp, 192	directed, 127
NetRule	

NetworkData, 126	Flock, 96
vertex_attr, 127	numbers
nfunctions	NetCounterData, 125
Flock, 98	,
Geese, 110	observed_counts
nfuns	Geese, 107
Flock, 95	offspring
Geese, 106	Node, 131
nleafs	ONE
Flock, 95	CHECK, 24
Geese, 107	EXISTS, 26
nnodes	operator Cell Type
Flock, 96	BArrayCell< Cell_Type, Data_Type >, 42
Geese, 107	BArrayCell_const< Cell_Type, Data_Type >, 44
nnozero	BArrayDenseCell< Cell_Type, Data_Type >, 62
BArray< Cell_Type, Data_Type >, 34	BArrayDenseCell_const< Cell_Type, Data_Type
BArrayDense < Cell_Type, Data_Type >, 54	>, 65
Node, 127	Cell< Cell_Type >, 77
	operator std::vector< Cell Type >
~Node, 129	BArrayVector< Cell_Type, Data_Type >, 69
annotations, 130	BArrayVector_const< Cell_Type, Data_Type >, 72
array, 130	operator!=
arrays, 131	BArrayCell_const< Cell_Type, Data_Type >, 44
duplication, 131	BArrayDenseCell_const< Cell_Type, Data_Type
get_parent, 130	>, 65
id, 131	BArrayVector_const< Cell_Type, Data_Type >, 72
is_leaf, 130	Cell< Cell_Type >, 77
narray, 131	operator<
Node, 128, 129	BArrayCell_const< Cell_Type, Data_Type >, 44
noffspring, 130	BArrayDenseCell_const< Cell_Type, Data_Type
offspring, 131	>, 65
ord, 132	BArrayVector_const< Cell_Type, Data_Type >, 73
parent, 132	operator<=
probability, 132	BArrayCell_const< Cell_Type, Data_Type >, 45
subtree_prob, 132 visited, 132	BArrayDenseCell_const< Cell_Type, Data_Type
NodeData, 133	>, 65
	BArrayVector_const< Cell_Type, Data_Type >, 73
∼NodeData, 134 blengths, 134	operator>
duplication, 134	BArrayCell_const< Cell_Type, Data_Type >, 45
•	BArrayDenseCell_const< Cell_Type, Data_Type
NodeData, 133, 134 states, 134	>, 66
nodes	BArrayVector_const< Cell_Type, Data_Type >, 73
Geese, 110	operator>=
noffspring	BArrayCell_const< Cell_Type, Data_Type >, 45
• •	BArrayDenseCell_const< Cell_Type, Data_Type
Node, 130 NONE	>, 66
	BArrayVector_const< Cell_Type, Data_Type >, 73
CHECK, 24	operator*=
EXISTS, 26	BArray< Cell_Type, Data_Type >, 35
NOW PArroy < Coll Type Date Type > 24	BArrayCell< Cell_Type, Data_Type >, 42
BArray< Cell_Type, Data_Type >, 34 BArrayDense< Cell_Type, Data_Type >, 54	BArrayDense< Cell_Type, Data_Type >, 55
	BArrayDenseCell< Cell_Type, Data_Type >, 62
nterms	BArray Vector < Cell_Type, Data_Type >, 69
Flock, 96	operator()
Geese, 107	BArray< Cell_Type, Data_Type >, 35
Model < Array_Type, Data_Counter_Type,	BArrayDense < Cell_Type, Data_Type >, 54
Data_Rule_Type, Data_Rule_Dyn_Type >,	Flock, 96
120	Rule< Array_Type, Data_Type >, 145
ntrees	Rules < Array_Type, Data_Type >, 148

vecHasher< T >, 163	counter_cogain, 16
operator+=	counter_gains, 16
BArray < Cell_Type, Data_Type >, 35	counter_gains_k_offspring, 17
BArrayCell< Cell_Type, Data_Type >, 42	
	counter_genes_changing, 17
BArrayDense < Cell_Type, Data_Type >, 55	counter_longest, 17
BArrayDenseCell< Cell_Type, Data_Type >, 63	counter_loss, 17
BArrayVector< Cell_Type, Data_Type >, 69	counter_maxfuns, 18
operator-=	counter_neofun, 18
BArray< Cell_Type, Data_Type >, 36	counter_neofun_a2b, 18
BArrayCell< Cell_Type, Data_Type >, 42	counter_overall_changes, 18
BArrayDense < Cell_Type, Data_Type >, 55, 56	counter_overall_gains, 19
BArrayDenseCell< Cell_Type, Data_Type >, 63	counter_overall_loss, 19
BArrayVector< Cell_Type, Data_Type >, 69	counter_subfun, 19
operator/=	Phylo rules, 20
BArray< Cell_Type, Data_Type >, 36	rule_dyn_limit_changes, 20
BArrayCell< Cell_Type, Data_Type >, 42	phylo.hpp
BArrayDense < Cell_Type, Data_Type >, 42	
	get_last_name, 198
BArrayDenseCell< Cell_Type, Data_Type >, 63	PHYLO_CHECK_MISSING, 195
BArrayVector< Cell_Type, Data_Type >, 69	PHYLO_COUNTER_LAMBDA, 195
operator=	PHYLO_RULE_DYN_LAMBDA, 196
BArray< Cell_Type, Data_Type >, 36	PhyloArray, 196
BArrayCell< Cell_Type, Data_Type >, 42	PhyloCounter, 196
BArrayDense < Cell_Type, Data_Type >, 56	PhyloCounterData, 196
BArrayDenseCell< Cell_Type, Data_Type >, 63	PhyloCounters, 197
BArrayVector< Cell_Type, Data_Type >, 70	PhyloModel, 197
Cell< Cell_Type >, 77, 78	PhyloPowerSet, 197
Counter< Array_Type, Data_Type >, 84	PhyloRule, 197
Counters< Array_Type, Data_Type >, 88, 89	PhyloRuleData, 197
	-
Geese, 107, 108	PhyloRuleDyn, 197
Model< Array_Type, Data_Counter_Type,	PhyloRules, 198
Data_Rule_Type, Data_Rule_Dyn_Type >,	PhyloRulesDyn, 198
120	PhyloStatsCounter, 198
Rules < Array_Type, Data_Type >, 148	PhyloSupport, 198
operator==	PHYLO_CHECK_MISSING
BArray< Cell_Type, Data_Type >, 37	phylo.hpp, 195
BArrayCell< Cell_Type, Data_Type >, 43	PHYLO_COUNTER_LAMBDA
BArrayCell_const< Cell_Type, Data_Type >, 45	phylo.hpp, 195
BArrayDense < Cell_Type, Data_Type >, 57	PHYLO_RULE_DYN_LAMBDA
BArrayDenseCell< Cell_Type, Data_Type >, 63	phylo.hpp, 196
BArrayDenseCell_const< Cell_Type, Data_Type	PhyloArray
	phylo.hpp, 196
>, 66	
BArrayVector< Cell_Type, Data_Type >, 70	PhyloCounter
BArrayVector_const< Cell_Type, Data_Type >, 73	phylo.hpp, 196
Cell< Cell_Type >, 78	PhyloCounterData
operator[]	phylo.hpp, 196
Counters< Array_Type, Data_Type >, 89	PhyloCounters
PowerSet< Array_Type, Data_Rule_Type >, 141	phylo.hpp, 197
ord	PhyloModel
Node, 132	phylo.hpp, 197
out_of_range	PhyloPowerSet
BArray< Cell_Type, Data_Type >, 37	phylo.hpp, 197
BArrayDense< Cell_Type, Data_Type >, 57	PhyloRule
DAITayDelise Cell_Type, Data_Type >, 37	-
parent	phylo.hpp, 197
Node, 132	PhyloRuleData
	phylo.hpp, 197
parse_polytomies	PhyloRuleDyn
Geese, 108	phylo.hpp, 197
Phylo counters, 15	PhyloRuleDynData, 135
counter_co_opt, 16	

\sim PhyloRuleDynData, 135	Data_Rule_Type, Data_Rule_Dyn_Type >
counts, 136	120
duplication, 136	printf_barry
lb, 136	barry-configuration.hpp, 180
PhyloRuleDynData, 135	probability
pos, 136	Node, 132
ub, 136	
PhyloRules	README.md, 224
phylo.hpp, 198	reduced_sequence
PhyloRulesDyn	Geese, 110
phylo.hpp, 198	rengine
PhyloStatsCounter	Flock, 98
phylo.hpp, 198	reserve
PhyloSupport	BArray< Cell_Type, Data_Type >, 37
phylo.hpp, 198	BArrayDense < Cell_Type, Data_Type >, 57
POS	FreqTable< T >, 100
barraydense-meet.hpp, 175	reset
• • • • • • • • • • • • • • • • • • • •	PowerSet< Array_Type, Data_Rule_Type >, 141
barraydensecell-meat.hpp, 177	reset_array
pos Divida Divida Divid	StatsCounter< Array_Type, Data_Type >, 151
PhyloRuleDynData, 136	Support< Array_Type, Data_Counter_Type,
PowerSet	Data_Rule_Type, Data_Rule_Dyn_Type >:
PowerSet< Array_Type, Data_Rule_Type >, 138	160
PowerSet < Array_Type, Data_Rule_Type >, 137	resize
\sim PowerSet, 139	
add_rule, 139	BArray Cell_Type, Data_Type >, 37
begin, 139	BArrayDense < Cell_Type, Data_Type >, 57
calc, 140	Entries< Cell_Type >, 91
coordinates_free, 141	rm_cell
coordinates_locked, 141	BArray< Cell_Type, Data_Type >, 37
data, 142	BArrayDense < Cell_Type, Data_Type >, 58
EmptyArray, 142	ROW
end, 140	barray-meat-operators.hpp, 168
get_data, 140	barray-meat.hpp, 170
get_data_ptr, 140	barraydense-meet.hpp, 175
init_support, 140	row
M, 142	BArray< Cell_Type, Data_Type >, 38
N, 142	BArrayDense < Cell_Type, Data_Type >, 58
operator[], 141	Row_type
PowerSet, 138	typedefs.hpp, 222
reset, 141	Rule
rules, 142	Rule < Array_Type, Data_Type >, 144
rules_deleted, 143	Rule < Array_Type, Data_Type >, 143
size, 141	~Rule, 144
•	D, 145
predict Coppe 108	operator(), 145
Geese, 108	Rule, 144
predict_backend	rule_dyn_limit_changes
Geese, 108	Phylo rules, 20
print	rule_fun_default
BArray < Cell_Type, Data_Type >, 37	rules-bones.hpp, 213
BArrayDense < Cell_Type, Data_Type >, 57	• •
FreqTable < T >, 100	Rule_fun_type
Support< Array_Type, Data_Counter_Type,	typedefs.hpp, 223
Data_Rule_Type, Data_Rule_Dyn_Type >,	RULE_FUNCTION
159	barry.hpp, 183
print_observed_counts	geese-bones.hpp, 204
Geese, 108	RULE_LAMBDA
print_stats	barry.hpp, 183
Model< Array_Type, Data_Counter_Type,	Rules
	Rules< Array_Type, Data_Type >, 146

rules	Geese, 109
PowerSet < Array_Type, Data_Rule_Type >, 142	Model < Array_Type, Data_Counter_Type,
Rules< Array_Type, Data_Type >, 145	Data_Rule_Type, Data_Rule_Dyn_Type >,
∼Rules, 146	122
add_rule, 147	simulate
clear, 147	Geese, 109
get_seq, 147	size
operator(), 148	BArrayVector< Cell_Type, Data_Type >, 70
operator=, 148	BArrayVector_const< Cell_Type, Data_Type >, 74
Rules, 146	Counters< Array_Type, Data_Type >, 90
size, 148	FreqTable< T >, 101
rules-bones.hpp	Model< Array_Type, Data_Counter_Type,
rule_fun_default, 213	Data_Rule_Type, Data_Rule_Dyn_Type >,
rules_deleted	122
PowerSet< Array_Type, Data_Rule_Type >, 143	PowerSet < Array_Type, Data_Rule_Type >, 141
rules_zerodiag	Rules < Array_Type, Data_Tune_Type >, 141
network.hpp, 193	size_unique
sample	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 >,	122
120, 121	source
	Entries < Cell_Type >, 92
sequence	states
Geese, 111	NodeData, 134
set_counters	Statistical Models, 9
Model < Array_Type, Data_Counter_Type,	StatsCounter
Data_Rule_Type, Data_Rule_Dyn_Type >,	StatsCounter< Array_Type, Data_Type >, 150
121	StatsCounter< Array_Type, Data_Type >, 149
StatsCounter< Array_Type, Data_Type >, 152	\sim StatsCounter, 150
Support< Array_Type, Data_Counter_Type,	add_counter, 150
Data_Rule_Type, Data_Rule_Dyn_Type >,	count_all, 151
160	count_current, 151
set_data	count_init, 151
BArray< Cell_Type, Data_Type >, 38	get_counters, 151
BArrayDense< Cell_Type, Data_Type >, 58	reset_array, 151
set_keygen	set_counters, 152
Model< Array_Type, Data_Counter_Type,	StatsCounter, 150
Data_Rule_Type, Data_Rule_Dyn_Type >,	store_psets
121	Model< Array_Type, Data_Counter_Type,
set_rengine	Data Rule Type, Data Rule Dyn Type >,
Model< Array_Type, Data_Counter_Type,	123
Data_Rule_Type, Data_Rule_Dyn_Type >,	subtree_prob
121	Node, 132
set_rules	Support
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 >,
122	155
Support< Array_Type, Data_Counter_Type,	Support < Array_Type, Data_Counter_Type, Data_Rule_Type
Data_Rule_Type, Data_Rule_Dyn_Type >,	Data_Rule_Dyn_Type >, 152
160	~Support, 156
set_rules_dyn	add_counter, 156
Model < Array_Type, Data_Counter_Type,	add_rule, 156, 157
Data_Rule_Type, Data_Rule_Dyn_Type >,	add_rule_dyn, 157
122	calc, 157
Support< Array_Type, Data_Counter_Type,	change_stats, 161
Data_Rule_Type, Data_Rule_Dyn_Type >,	coordinates_free, 161
160	coordinates_free, 161
set seed	
Flock, 97	current_stats, 161
•	

delete_counters, 162	uint, 223
delete rules, 162	vec_equal, 223
delete_rules_dyn, 162	vec_equal_approx, 223
get_counters, 158	vec_inner_prod, 224
get_counts, 158	
get_counts_ptr, 158	ub
get_current_stats, 158	PhyloRuleDynData, 136
-	uint
get_data, 158	typedefs.hpp, 223
get_rules, 159	UKNOWN
get_rules_dyn, 159	EXISTS, 26
init_support, 159	
M, 162	update_annotations
max_num_elements, 162	Geese, 109
N, 163	update_normalizing_constant
print, 159	model-bones.hpp, 200
reset_array, 160	
set_counters, 160	val
set_rules, 160	Entries < Cell_Type >, 92
set_rules_dyn, 160	value
Support, 155	Cell< Cell_Type >, 78
support-meat.hpp	vec_diff
BARRY_SUPPORT_MEAT_HPP, 220	geese-bones.hpp, 204
support_size	vec_equal
Flock, 97	typedefs.hpp, 223
Geese, 109	vec_equal_approx
	typedefs.hpp, 223
3=31 / = = 31 /	vec_inner_prod
Data_Rule_Type, Data_Rule_Dyn_Type >,	typedefs.hpp, 224
123	vecHasher< T >, 163
swap_cells	operator(), 163
BArray< Cell_Type, Data_Type >, 38	vector_caster
BArrayDense < Cell_Type, Data_Type >, 59	geese-bones.hpp, 204
swap_cols	
BArray< Cell_Type, Data_Type >, 38	vertex_attr
BArrayDense< Cell_Type, Data_Type >, 59	NetworkData, 127
swap_rows	visited
BArray< Cell_Type, Data_Type >, 39	BArray < Cell_Type, Data_Type >, 40
BArrayDense < Cell_Type, Data_Type >, 59	BArrayDense < Cell_Type, Data_Type >, 61
	Cell< Cell_Type >, 78
target	Node, 132
Entries< Cell_Type >, 92	7570 0511
toggle_cell	ZERO_CELL
BArray< Cell_Type, Data_Type >, 39	barraydense-meet.hpp, 175
BArrayDense < Cell_Type, Data_Type >, 59	zero_col
toggle_lock	BArray< Cell_Type, Data_Type >, 39
BArray< Cell_Type, Data_Type >, 39	BArrayDense < Cell_Type, Data_Type >, 60
BArrayDense< Cell_Type, Data_Type >, 59	zero_row
transpose	BArray< Cell_Type, Data_Type >, 40
BArray< Cell_Type, Data_Type >, 39	BArrayDense < Cell_Type, Data_Type >, 60
BArrayDense < Cell_Type, Data_Type >, 60	_, _, _,
TWO	
CHECK, 24	
EXISTS, 26	
typedefs.hpp	
Col_type, 222	
Counter_fun_type, 222	
Counts_type, 222	
MapVec_type, 222	
Row_type, 222	
Rule_fun_type, 223	