barry: Your go-to motif accountant 0.0-1

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1 Main Page	1
2 Module Index	5
2.1 Modules	5
3 Class Index	7
3.1 Class List	7
4 File Index	9
4.1 File List	9
5 Module Documentation	11
5.1 Counting	11
5.1.1 Detailed Description	11
5.2 Statistical Models	11
5.2.1 Detailed Description	12
5.3 Network counters	12
5.3.1 Detailed Description	13
5.3.2 Function Documentation	13
5.3.2.1 counter_absdiff()	13
5.3.2.2 counter_ctriads() [1/2]	14
5.3.2.3 counter_ctriads() [2/2]	14
5.3.2.4 counter_degree()	14
5.3.2.5 counter_density()	14
5.3.2.6 counter_diff()	14
5.3.2.7 counter_edges()	15
5.3.2.8 counter_idegree() [1/2]	15
5.3.2.9 counter_idegree() [2/2]	15
5.3.2.10 counter_idegree15() [1/2]	15
5.3.2.11 counter idegree15() [2/2]	15
5.3.2.12 counter_isolates() [1/2]	16
5.3.2.13 counter_isolates() [2/2]	16
5.3.2.14 counter istar2() [1/2]	16
5.3.2.15 counter_istar2() [2/2]	16
5.3.2.16 counter_mutual()	16
5.3.2.17 counter_nodecov()	17
5.3.2.18 counter_nodeicov()	17
5.3.2.19 counter_nodematch()	17
5.3.2.20 counter_nodeocov()	17
5.3.2.21 counter_odegree() [1/2]	17
5.3.2.22 counter_odegree() [2/2]	18
5.3.2.23 counter_odegree15() [1/2]	18
5.3.2.24 counter_odegree15() [2/2]	18
5.3.2.25 counter_ostar2() [1/2]	18
0.0.E.20 00ditol_00tdt2([1/2]	10

5.3.2.26 counter_ostar2() [2/2]	18
5.3.2.27 counter_ttriads() [1/2]	19
5.3.2.28 counter_ttriads() [2/2]	19
5.3.2.29 NETWORK_COUNTER()	19
5.4 Phylo counters	19
5.4.1 Detailed Description	20
5.4.2 Function Documentation	21
5.4.2.1 counter_co_opt()	21
5.4.2.2 counter_cogain()	21
5.4.2.3 counter_gains()	21
5.4.2.4 counter_gains_from_0()	22
5.4.2.5 counter_gains_k_offspring()	22
5.4.2.6 counter_genes_changing()	22
5.4.2.7 counter_k_genes_changing()	22
5.4.2.8 counter_less_than_p_prop_genes_changing()	23
5.4.2.9 counter_longest()	23
5.4.2.10 counter_loss()	23
5.4.2.11 counter_maxfuns()	23
5.4.2.12 counter_neofun()	24
5.4.2.13 counter_neofun_a2b()	24
5.4.2.14 counter_overall_changes()	24
5.4.2.15 counter_overall_gains()	24
5.4.2.16 counter_overall_gains_from_0()	25
5.4.2.17 counter_overall_loss()	25
5.4.2.18 counter_pairwise_first_gain()	25
5.4.2.19 counter_pairwise_neofun_singlefun()	25
5.4.2.20 counter_pairwise_overall_change()	26
5.4.2.21 counter_pairwise_preserving()	26
5.4.2.22 counter_preserve_pseudogene()	26
5.4.2.23 counter_prop_genes_changing()	26
5.4.2.24 counter_subfun()	27
5.5 Phylo rules	27
5.5.1 Detailed Description	27
5.5.2 Function Documentation	27
5.5.2.1 rule_dyn_limit_changes()	27
6 Namespace Documentation	29
6.1 barry Namespace Reference	29
6.1.1 Detailed Description	29
6.2 barry::counters Namespace Reference	29
6.2.1 Detailed Description	29
6.3 barry::counters::network Namespace Reference	30

6.4 barry::counters::phylo Namespace Reference	 30
6.5 CHECK Namespace Reference	 30
6.5.1 Detailed Description	 30
6.5.2 Variable Documentation	 30
6.5.2.1 BOTH	 30
6.5.2.2 NONE	 30
6.5.2.3 ONE	 30
6.5.2.4 TWO	 31
6.6 EXISTS Namespace Reference	 31
6.6.1 Detailed Description	 31
6.6.2 Variable Documentation	 31
6.6.2.1 AS_ONE	 31
6.6.2.2 AS_ZERO	 31
6.6.2.3 BOTH	 32
6.6.2.4 NONE	 32
6.6.2.5 ONE	 32
6.6.2.6 TWO	 32
6.6.2.7 UKNOWN	 32
7 Class Documentation	33
7.1 BArray < Cell_Type, Data_Type > Class Template Reference	33
7.1 BATTay Cell_Type, Data_Type > Class Template Reference	35
7.1.2 Constructor & Destructor Documentation	36
7.1.2 Constructor & Destructor Documentation	36
7.1.2.1 BAITay() [1/6]	36
7.1.2.2 BArray() [2/6]	36
7.1.2.4 BArray() [4/6]	37
7.1.2.5 BArray() [5/6]	37
7.1.2.6 BArray() [6/6]	37
7.1.2.7 ~BArray()	37
7.1.3 Member Function Documentation	37
7.1.3.1 clear()	37
7.1.3.2 col()	38
7.1.3.3 D() [1/2]	38
7.1.3.4 D() [2/2]	38
7.1.3.5 default_val()	38
7.1.3.6 flush_data()	38
7.1.3.7 get_cell()	38
7.1.3.8 get_col_vec() [1/2]	39
7.1.3.9 get_col_vec() [2/2]	39
7.1.3.10 get_entries()	39
7.1.3.11 get_row_vec() [1/2]	39
	_

7.1.3.12 get_row_vec() [2/2]	38
7.1.3.13 insert_cell() [1/3]	40
7.1.3.14 insert_cell() [2/3]	40
7.1.3.15 insert_cell() [3/3]	40
7.1.3.16 is_dense()	40
7.1.3.17 is_empty()	40
7.1.3.18 ncol()	41
7.1.3.19 nnozero()	41
7.1.3.20 nrow()	41
7.1.3.21 operator()() [1/2]	41
7.1.3.22 operator()() [2/2]	41
7.1.3.23 operator*=()	41
7.1.3.24 operator+=() [1/3]	42
7.1.3.25 operator+=() [2/3]	42
7.1.3.26 operator+=() [3/3]	42
7.1.3.27 operator-=() [1/3]	42
7.1.3.28 operator-=() [2/3]	42
7.1.3.29 operator-=() [3/3]	42
7.1.3.30 operator/=()	43
7.1.3.31 operator=() [1/2]	43
7.1.3.32 operator=() [2/2]	43
7.1.3.33 operator==()	43
7.1.3.34 out_of_range()	43
7.1.3.35 print()	43
7.1.3.36 reserve()	44
7.1.3.37 resize()	44
7.1.3.38 rm_cell()	44
7.1.3.39 row()	44
7.1.3.40 set_data()	
7.1.3.41 swap_cells()	45
7.1.3.42 swap_cols()	45
7.1.3.43 swap_rows()	45
7.1.3.44 toggle_cell()	45
7.1.3.45 toggle_lock()	46
7.1.3.46 transpose()	46
7.1.3.47 zero_col()	46
7.1.3.48 zero_row()	46
7.1.4 Friends And Related Function Documentation	46
7.1.4.1 BArrayCell < Cell_Type, Data_Type >	46
7.1.4.2 BArrayCell_const< Cell_Type, Data_Type >	47
7.1.5 Member Data Documentation	47
7.1.5.1 visited	47

7.2 BArrayCell< Cell_Type, Data_Type > Class Template Reference	47
7.2.1 Detailed Description	47
7.2.2 Constructor & Destructor Documentation	48
7.2.2.1 BArrayCell()	48
7.2.2.2 ~BArrayCell()	48
7.2.3 Member Function Documentation	48
7.2.3.1 operator Cell_Type()	48
7.2.3.2 operator*=()	48
7.2.3.3 operator+=()	49
7.2.3.4 operator-=()	49
7.2.3.5 operator/=()	49
7.2.3.6 operator=()	49
7.2.3.7 operator==()	49
7.3 BArrayCell_const< Cell_Type, Data_Type > Class Template Reference	50
7.3.1 Detailed Description	50
7.3.2 Constructor & Destructor Documentation	50
7.3.2.1 BArrayCell_const()	50
7.3.2.2 ~BArrayCell_const()	50
7.3.3 Member Function Documentation	51
7.3.3.1 operator Cell_Type()	51
7.3.3.2 operator"!=()	51
7.3.3.3 operator<()	51
7.3.3.4 operator<=()	51
7.3.3.5 operator==()	51
7.3.3.6 operator>()	52
7.3.3.7 operator>=()	52
7.4 BArrayDense< Cell_Type, Data_Type > Class Template Reference	52
7.4.1 Detailed Description	54
7.4.2 Constructor & Destructor Documentation	55
7.4.2.1 BArrayDense() [1/6]	55
7.4.2.2 BArrayDense() [2/6]	55
7.4.2.3 BArrayDense() [3/6]	55
7.4.2.4 BArrayDense() [4/6]	56
7.4.2.5 BArrayDense() [5/6]	56
7.4.2.6 BArrayDense() [6/6]	56
7.4.2.7 ~BArrayDense()	56
7.4.3 Member Function Documentation	56
7.4.3.1 clear()	56
7.4.3.2 col() [1/2]	57
7.4.3.3 col() [2/2]	57
7.4.3.4 colsum()	57
7.4.3.5 D() [1/2]	57

7.4.3.6 D() [2/2]
7.4.3.7 default_val()
7.4.3.8 get_cell()
7.4.3.9 get_col_vec() [1/2] 58
7.4.3.10 get_col_vec() [2/2]
7.4.3.11 get_data()
7.4.3.12 get_entries()
7.4.3.13 get_row_vec() [1/2] 59
7.4.3.14 get_row_vec() [2/2]
7.4.3.15 insert_cell() [1/2]
7.4.3.16 insert_cell() [2/2]
7.4.3.17 is_dense()
7.4.3.18 is_empty()
7.4.3.19 ncol()
7.4.3.20 nnozero()
7.4.3.21 nrow()
7.4.3.22 operator()() [1/2]
7.4.3.23 operator()() [2/2]
7.4.3.24 operator*=()
7.4.3.25 operator+=() [1/3]
7.4.3.26 operator+=() [2/3]
7.4.3.27 operator+=() [3/3]
7.4.3.28 operator-=() [1/3]
7.4.3.29 operator-=() [2/3]
7.4.3.30 operator-=() [3/3]
7.4.3.31 operator/=()
7.4.3.32 operator=() [1/2]
7.4.3.33 operator=() [2/2]
7.4.3.34 operator==()
7.4.3.35 out_of_range()
7.4.3.36 print()
7.4.3.37 reserve()
7.4.3.38 resize()
7.4.3.39 rm_cell()
7.4.3.40 row() [1/2]
7.4.3.41 row() [2/2]
7.4.3.42 rowsum()
7.4.3.43 set_data()
7.4.3.44 swap_cells()
7.4.3.45 swap_cols()
7.4.3.46 swap_rows()
7.4.3.47 toggle_cell()

7.4.3.48 toggle_lock()	65
7.4.3.49 transpose()	65
7.4.3.50 zero_col()	65
7.4.3.51 zero_row()	66
7.4.4 Friends And Related Function Documentation	66
7.4.4.1 BArrayDenseCell< Cell_Type, Data_Type >	66
7.4.4.2 BArrayDenseCol< Cell_Type, Data_Type >	66
7.4.4.3 BArrayDenseCol_const< Cell_Type, Data_Type >	66
7.4.4.4 BArrayDenseRow < Cell_Type, Data_Type >	66
7.4.4.5 BArrayDenseRow_const< Cell_Type, Data_Type >	67
7.4.5 Member Data Documentation	67
7.4.5.1 visited	67
7.5 BArrayDenseCell< Cell_Type, Data_Type > Class Template Reference	67
7.5.1 Detailed Description	68
7.5.2 Constructor & Destructor Documentation	68
7.5.2.1 BArrayDenseCell()	68
7.5.2.2 ~BArrayDenseCell()	68
7.5.3 Member Function Documentation	68
7.5.3.1 operator Cell_Type()	68
7.5.3.2 operator*=()	69
7.5.3.3 operator+=()	69
7.5.3.4 operator-=()	69
7.5.3.5 operator/=()	69
7.5.3.6 operator=()	69
7.5.3.7 operator==()	70
7.5.4 Friends And Related Function Documentation	70
7.5.4.1 BArrayDense < Cell_Type, Data_Type >	70
7.5.4.2 BArrayDenseCol< Cell_Type, Data_Type >	70
7.5.4.3 BArrayDenseCol_const< Cell_Type, Data_Type >	70
7.6 BArrayDenseCell_const< Cell_Type, Data_Type > Class Template Reference	71
7.6.1 Detailed Description	71
7.7 BArrayDenseCol< Cell_Type, Data_Type > Class Template Reference	71
7.7.1 Detailed Description	71
7.7.2 Constructor & Destructor Documentation	71
7.7.2.1 BArrayDenseCol()	72
7.7.3 Member Function Documentation	72
7.7.3.1 begin()	72
7.7.3.2 end()	72
7.7.3.3 operator()()	72
7.7.3.4 size()	72
7.7.4 Friends And Related Function Documentation	73
7.7.4.1 BArrayDense< Cell Type, Data Type >	73

7.7.4.2 BArrayDenseCell< Cell_Type, Data_Type >	73
7.7.4.3 BArrayDenseCell_const< Cell_Type, Data_Type >	73
7.8 BArrayDenseCol_const< Cell_Type, Data_Type > Class Template Reference	73
7.8.1 Detailed Description	74
7.8.2 Constructor & Destructor Documentation	74
7.8.2.1 BArrayDenseCol_const()	74
7.8.3 Member Function Documentation	74
7.8.3.1 begin()	74
7.8.3.2 end()	74
7.8.3.3 operator()()	75
7.8.3.4 size()	75
7.8.4 Friends And Related Function Documentation	75
7.8.4.1 BArrayDenseCell< Cell_Type, Data_Type >	75
7.8.4.2 BArrayDenseCell_const< Cell_Type, Data_Type >	75
7.9 BArrayDenseRow< Cell_Type, Data_Type > Class Template Reference	75
7.9.1 Detailed Description	76
7.9.2 Constructor & Destructor Documentation	76
7.9.2.1 BArrayDenseRow()	76
7.9.3 Member Function Documentation	76
7.9.3.1 begin()	76
7.9.3.2 end()	77
7.9.3.3 operator()()	77
7.9.3.4 size()	77
7.9.4 Friends And Related Function Documentation	77
7.9.4.1 BArrayDense < Cell_Type, Data_Type >	77
7.9.4.2 BArrayDenseCell< Cell_Type, Data_Type >	77
7.9.4.3 BArrayDenseCell_const< Cell_Type, Data_Type >	78
7.10 BArrayDenseRow_const< Cell_Type, Data_Type > Class Template Reference	78
7.10.1 Detailed Description	78
7.10.2 Constructor & Destructor Documentation	78
7.10.2.1 BArrayDenseRow_const()	79
7.10.3 Member Function Documentation	79
7.10.3.1 begin()	79
7.10.3.2 end()	79
7.10.3.3 operator()()	79
7.10.3.4 size()	79
7.10.4 Friends And Related Function Documentation	80
7.10.4.1 BArrayDenseCell< Cell_Type, Data_Type >	80
7.10.4.2 BArrayDenseCell_const< Cell_Type, Data_Type >	80
7.11 BArrayRow< Cell_Type, Data_Type > Class Template Reference	80
7.11.1 Detailed Description	80
7.11.2 Constructor & Destructor Documentation	81

7.11.2.1 BArrayRow()	81
7.11.2.2 ~BArrayRow()	81
7.11.3 Member Function Documentation	81
7.11.3.1 operator BArrayRow< Cell_Type, Data_Type >()	81
7.11.3.2 operator*=()	81
7.11.3.3 operator+=()	81
7.11.3.4 operator-=()	82
7.11.3.5 operator/=()	82
7.11.3.6 operator=()	82
7.11.3.7 operator==()	82
7.12 BArrayRow_const< Cell_Type, Data_Type > Class Template Reference	82
7.12.1 Detailed Description	83
7.12.2 Constructor & Destructor Documentation	83
7.12.2.1 BArrayRow_const()	83
7.12.2.2 ~BArrayRow_const()	83
7.12.3 Member Function Documentation	83
7.12.3.1 operator BArrayRow_const< Cell_Type, Data_Type >()	83
7.12.3.2 operator"!=()	83
7.12.3.3 operator<()	84
7.12.3.4 operator<=()	84
7.12.3.5 operator==()	84
7.12.3.6 operator>()	84
7.12.3.7 operator>=()	84
7.13 BArrayVector< Cell_Type, Data_Type > Class Template Reference	84
7.13.1 Detailed Description	85
7.13.2 Constructor & Destructor Documentation	85
7.13.2.1 BArrayVector()	85
7.13.2.2 ~BArrayVector()	86
7.13.3 Member Function Documentation	86
7.13.3.1 begin()	86
7.13.3.2 end()	86
7.13.3.3 is_col()	86
7.13.3.4 is_row()	87
7.13.3.5 operator std::vector< Cell_Type >()	87
7.13.3.6 operator*=()	87
7.13.3.7 operator+=()	87
7.13.3.8 operator-=()	87
7.13.3.9 operator/=()	88
7.13.3.10 operator=()	88
7.13.3.11 operator==()	88
7.13.3.12 size()	88
7.14 BArrayVector_const< Cell_Type, Data_Type > Class Template Reference	88

7.14.1 Detailed Description	89
7.14.2 Constructor & Destructor Documentation	89
7.14.2.1 BArrayVector_const()	89
7.14.2.2 ~BArrayVector_const()	89
7.14.3 Member Function Documentation	90
7.14.3.1 begin()	90
7.14.3.2 end()	90
7.14.3.3 is_col()	90
7.14.3.4 is_row()	90
7.14.3.5 operator std::vector< Cell_Type >()	90
7.14.3.6 operator"!=()	91
7.14.3.7 operator<()	91
7.14.3.8 operator<=()	91
7.14.3.9 operator==()	91
7.14.3.10 operator>()	91
7.14.3.11 operator>=()	92
7.14.3.12 size()	92
7.15 Cell < Cell_Type > Class Template Reference	92
7.15.1 Detailed Description	93
7.15.2 Constructor & Destructor Documentation	93
7.15.2.1 Cell() [1/7]	93
7.15.2.2 Cell() [2/7]	93
7.15.2.3 ~Cell()	93
7.15.2.4 Cell() [3/7]	94
7.15.2.5 Cell() [4/7]	94
7.15.2.6 Cell() [5/7]	94
7.15.2.7 Cell() [6/7]	94
7.15.2.8 Cell() [7/7]	94
7.15.3 Member Function Documentation	94
7.15.3.1 add() [1/4]	95
7.15.3.2 add() [2/4]	95
7.15.3.3 add() [3/4]	95
7.15.3.4 add() [4/4]	95
7.15.3.5 operator Cell_Type()	95
7.15.3.6 operator"!=()	95
7.15.3.7 operator=() [1/2]	96
7.15.3.8 operator=() [2/2]	96
7.15.3.9 operator==()	96
7.15.4 Member Data Documentation	96
7.15.4.1 active	96
7.15.4.2 value	96
7.15.4.3 visited	97

7.16 Cell_const< Cell_Type > Class Template Reference	97
7.16.1 Detailed Description	97
7.17 ConstBArrayRowlter< Cell_Type, Data_Type > Class Template Reference	97
7.17.1 Detailed Description	98
7.17.2 Constructor & Destructor Documentation	98
7.17.2.1 ConstBArrayRowlter()	98
7.17.2.2 ~ConstBArrayRowlter()	98
7.17.3 Member Data Documentation	98
7.17.3.1 Array	99
7.17.3.2 current_col	99
7.17.3.3 current_row	99
7.17.3.4 iter	99
7.18 Counter< Array_Type, Data_Type > Class Template Reference	99
7.18.1 Detailed Description)0
7.18.2 Constructor & Destructor Documentation)0
7.18.2.1 Counter() [1/4])1
7.18.2.2 Counter() [2/4])1
7.18.2.3 Counter() [3/4])1
7.18.2.4 Counter() [4/4])1
7.18.2.5 ~Counter())1
7.18.3 Member Function Documentation)2
7.18.3.1 count())2
7.18.3.2 get_description())2
7.18.3.3 get_name())2
7.18.3.4 init())2
7.18.3.5 operator=() [1/2])2
7.18.3.6 operator=() [2/2])3
7.18.4 Member Data Documentation)3
7.18.4.1 count_fun)3
7.18.4.2 data)3
7.18.4.3 desc)3
7.18.4.4 init_fun)3
7.18.4.5 name)4
7.19 Counters < Array_Type, Data_Type > Class Template Reference)4
7.19.1 Detailed Description)4
7.19.2 Constructor & Destructor Documentation)5
7.19.2.1 Counters() [1/3])5
7.19.2.2 ~Counters())5
7.19.2.3 Counters() [2/3])5
7.19.2.4 Counters() [3/3])5
7.19.3 Member Function Documentation)6
7.19.3.1 add_counter() [1/2])6

7.19.3.2 add_counter() [2/2]	 106
7.19.3.3 get_descriptions()	 106
7.19.3.4 get_names()	 106
7.19.3.5 operator=() [1/2]	 106
7.19.3.6 operator=() [2/2]	 107
7.19.3.7 operator[]()	 107
7.19.3.8 size()	 108
7.20 Entries < Cell_Type > Class Template Reference	 108
7.20.1 Detailed Description	 108
7.20.2 Constructor & Destructor Documentation	 109
7.20.2.1 Entries() [1/2]	 109
7.20.2.2 Entries() [2/2]	 109
7.20.2.3 ~Entries()	 109
7.20.3 Member Function Documentation	 109
7.20.3.1 resize()	 109
7.20.4 Member Data Documentation	 109
7.20.4.1 source	 110
7.20.4.2 target	 110
7.20.4.3 val	 110
7.21 Flock Class Reference	 110
7.21.1 Detailed Description	 111
7.21.2 Constructor & Destructor Documentation	 112
7.21.2.1 Flock()	 112
7.21.2.2 ~Flock()	 112
7.21.3 Member Function Documentation	 112
7.21.3.1 add_data()	 112
7.21.3.2 colnames()	 113
7.21.3.3 get_counters()	 113
7.21.3.4 get_model()	 113
7.21.3.5 get_stats_support()	 113
7.21.3.6 get_stats_target()	 113
7.21.3.7 get_support_fun()	 113
7.21.3.8 init()	 114
7.21.3.9 likelihood_joint()	 114
7.21.3.10 nfuns()	 114
7.21.3.11 nleafs()	 114
7.21.3.12 nnodes()	 115
7.21.3.13 nterms()	 115
7.21.3.14 ntrees()	 115
7.21.3.15 operator()()	 115
7.21.3.16 parse_polytomies()	 115
7.21.3.17 print()	 116

7.21.3.18 set_seed()	116
7.21.3.19 support_size()	116
7.21.4 Member Data Documentation	
7.21.4.1 dat	116
7.21.4.2 initialized	117
7.21.4.3 model	117
7.21.4.4 nfunctions	117
7.21.4.5 rengine	117
7.22 FreqTable < T > Class Template Reference	117
7.22.1 Detailed Description	118
7.22.2 Constructor & Destructor Documentation	118
7.22.2.1 FreqTable()	118
7.22.2.2 ∼FreqTable()	119
7.22.3 Member Function Documentation	119
7.22.3.1 add()	119
7.22.3.2 as_vector()	119
7.22.3.3 clear()	119
7.22.3.4 get_data()	119
7.22.3.5 get_index()	120
7.22.3.6 make_hash()	120
7.22.3.7 print()	120
7.22.3.8 reserve()	120
7.22.3.9 size()	120
7.23 Geese Class Reference	121
7.23.1 Detailed Description	124
7.23.2 Constructor & Destructor Documentation	124
7.23.2.1 Geese() [1/8]	124
7.23.2.2 Geese() [2/8]	125
7.23.2.3 Geese() [3/8]	125
7.23.2.4 Geese() [4/8]	125
7.23.2.5 ~Geese() [1/2]	125
7.23.2.6 Geese() [5/8]	125
7.23.2.7 Geese() [6/8]	126
7.23.2.8 Geese() [7/8]	126
7.23.2.9 Geese() [8/8]	126
7.23.2.10 ~Geese() [2/2]	126
7.23.3 Member Function Documentation	126
7.23.3.1 calc_reduced_sequence() [1/2]	126
7.23.3.2 calc_reduced_sequence() [2/2]	126
7.23.3.3 calc_sequence() [1/2]	127
7.23.3.4 calc_sequence() [2/2]	127
7.23.3.5 colnames() [1/2]	127

7.23.3.6 colnames() [2/2]
7.23.3.7 get_annotated_nodes() [1/2]
7.23.3.8 get_annotated_nodes() [2/2]
7.23.3.9 get_counters() [1/2]
7.23.3.10 get_counters() [2/2]
7.23.3.11 get_model() [1/2]
7.23.3.12 get_model() [2/2]
7.23.3.13 get_probabilities() [1/2]
7.23.3.14 get_probabilities() [2/2]
7.23.3.15 get_rengine() [1/2]
7.23.3.16 get_rengine() [2/2]
7.23.3.17 get_states() [1/2]
7.23.3.18 get_states() [2/2]
7.23.3.19 get_support_fun() [1/2]
7.23.3.20 get_support_fun() [2/2]
7.23.3.21 inherit_support() [1/2]
7.23.3.22 inherit_support() [2/2]
7.23.3.23 init() [1/2]
7.23.3.24 init() [2/2]
7.23.3.25 init_node() [1/2]
7.23.3.26 init_node() [2/2]
7.23.3.27 likelihood() [1/2]
7.23.3.28 likelihood() [2/2]
7.23.3.29 likelihood_exhaust() [1/2]
7.23.3.30 likelihood_exhaust() [2/2]
7.23.3.31 nannotations() [1/2]
7.23.3.32 nannotations() [2/2]
7.23.3.33 nfuns() [1/2]
7.23.3.34 nfuns() [2/2]
7.23.3.35 nleafs() [1/2]
7.23.3.36 nleafs() [2/2]
7.23.3.37 nnodes() [1/2]
7.23.3.38 nnodes() [2/2]
7.23.3.39 nterms() [1/2]
7.23.3.40 nterms() [2/2]
7.23.3.41 observed_counts() [1/2]
7.23.3.42 observed_counts() [2/2]
7.23.3.43 operator=() [1/4]
7.23.3.44 operator=() [2/4]
7.23.3.45 operator=() [3/4]
7.23.3.46 operator=() [4/4]
7.23.3.47 parse_polytomies() [1/2]

7.23.3.48 parse_polytomies() [2/2]	134
7.23.3.49 predict() [1/2]	134
7.23.3.50 predict() [2/2]	135
7.23.3.51 predict_backend() [1/2]	135
7.23.3.52 predict_backend() [2/2]	135
7.23.3.53 predict_exhaust() [1/2]	135
7.23.3.54 predict_exhaust() [2/2]	135
7.23.3.55 predict_exhaust_backend() [1/2]	136
7.23.3.56 predict_exhaust_backend() [2/2]	136
7.23.3.57 predict_sim() [1/2]	136
7.23.3.58 predict_sim() [2/2]	136
7.23.3.59 print() [1/2]	136
7.23.3.60 print() [2/2]	137
7.23.3.61 print_observed_counts() [1/2]	137
7.23.3.62 print_observed_counts() [2/2]	137
7.23.3.63 set_seed() [1/2]	137
7.23.3.64 set_seed() [2/2]	137
7.23.3.65 simulate() [1/2]	137
7.23.3.66 simulate() [2/2]	138
7.23.3.67 support_size() [1/2]	138
7.23.3.68 support_size() [2/2]	138
7.23.3.69 update_annotations() [1/2]	138
7.23.3.70 update_annotations() [2/2]	138
7.23.4 Member Data Documentation	138
7.23.4.1 delete_rengine	139
7.23.4.2 delete_support	139
7.23.4.3 initialized	139
7.23.4.4 map_to_nodes	139
7.23.4.5 nfunctions	139
7.23.4.6 nodes	139
7.23.4.7 pset_loc	140
7.23.4.8 reduced_sequence	
7.23.4.9 sequence	140
7.24 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class Templete Performance	4.40
plate Reference	
7.24.1 Detailed Description	
7.24.2 Constructor & Destructor Documentation	
7.24.2.1 Model() [1/3]	
7.24.2.2 Model() [2/3]	
7.24.2.3 Model() [3/3]	
7.24.2.4 ~ Model()	
7.24.3 Member Function Documentation	144

7.24.3.1 add_array()
7.24.3.2 add_counter() [1/2]
7.24.3.3 add_counter() [2/2]
7.24.3.4 add_rule() [1/2]
7.24.3.5 add_rule() [2/2]
7.24.3.6 add_rule_dyn() [1/2]
7.24.3.7 add_rule_dyn() [2/2]
7.24.3.8 colnames()
7.24.3.9 conditional_prob()
7.24.3.10 gen_key()
7.24.3.11 get_arrays2support()
7.24.3.12 get_counters()
7.24.3.13 get_norm_const()
7.24.3.14 get_pset()
7.24.3.15 get_pset_arrays()
7.24.3.16 get_pset_probs()
7.24.3.17 get_pset_stats() [1/2]
7.24.3.18 get_pset_stats() [2/2]
7.24.3.19 get_rengine()
7.24.3.20 get_rules()
7.24.3.21 get_rules_dyn()
7.24.3.22 get_stats_support()
7.24.3.23 get_stats_target()
7.24.3.24 get_support_fun()
7.24.3.25 likelihood() [1/3]
7.24.3.26 likelihood() [2/3]
7.24.3.27 likelihood() [3/3]
7.24.3.28 likelihood_total()
7.24.3.29 nterms()
7.24.3.30 operator=()
7.24.3.31 print()
7.24.3.32 print_stats()
7.24.3.33 sample() [1/2]
7.24.3.34 sample() [2/2]
7.24.3.35 set_counters()
7.24.3.36 set_keygen()
7.24.3.37 set_rengine()
7.24.3.38 set_rules()
7.24.3.39 set_rules_dyn()
7.24.3.40 set_seed()
7.24.3.41 set_transform_model()
7.24.3.42 size()

7.24.3.43 size_unique()	54
7.24.3.44 store_psets()	54
7.24.3.45 support_size()	54
7.24.3.46 transform_model()	54
7.25 NetCounterData Class Reference	54
7.25.1 Detailed Description	55
7.25.2 Constructor & Destructor Documentation	55
7.25.2.1 NetCounterData() [1/2]	55
7.25.2.2 NetCounterData() [2/2]	55
7.25.2.3 ~NetCounterData()	55
7.25.3 Member Data Documentation	56
7.25.3.1 indices	56
7.25.3.2 numbers	56
7.26 NetworkData Class Reference	56
7.26.1 Detailed Description	57
7.26.2 Constructor & Destructor Documentation	57
7.26.2.1 NetworkData() [1/3]	57
7.26.2.2 NetworkData() [2/3]	57
7.26.2.3 NetworkData() [3/3]	57
7.26.2.4 ∼NetworkData()	58
7.26.3 Member Data Documentation	58
7.26.3.1 directed	58
7.26.3.2 vertex_attr	58
7.27 Node Class Reference	58
7.27.1 Detailed Description	59
7.27.2 Constructor & Destructor Documentation	59
7.27.2.1 Node() [1/5]	30
7.27.2.2 Node() [2/5]	30
7.27.2.3 Node() [3/5]	30
7.27.2.4 Node() [4/5]	30
7.27.2.5 Node() [5/5]	30
7.27.2.6 ∼Node()	31
7.27.3 Member Function Documentation	31
7.27.3.1 get_parent()	31
7.27.3.2 is_leaf()	31
7.27.3.3 noffspring()	31
7.27.4 Member Data Documentation	31
7.27.4.1 annotations	31
7.27.4.2 array	32
7.27.4.3 arrays	32
7.27.4.4 duplication	32
7.27.4.5 id	32

7.27.4.6 narray	162
7.27.4.7 offspring	163
7.27.4.8 ord	163
7.27.4.9 parent	163
7.27.4.10 probability	163
7.27.4.11 subtree_prob	163
7.27.4.12 visited	164
7.28 NodeData Class Reference	164
7.28.1 Detailed Description	164
7.28.2 Constructor & Destructor Documentation	164
7.28.2.1 NodeData()	164
7.28.3 Member Data Documentation	165
7.28.3.1 blengths	165
7.28.3.2 duplication	165
7.28.3.3 states	165
7.29 PhyloCounterData Class Reference	165
7.29.1 Detailed Description	166
7.29.2 Constructor & Destructor Documentation	166
7.29.2.1 PhyloCounterData() [1/2]	166
7.29.2.2 PhyloCounterData() [2/2]	166
7.29.3 Member Function Documentation	166
7.29.3.1 at()	166
7.29.3.2 begin()	166
7.29.3.3 empty()	167
7.29.3.4 end()	167
7.29.3.5 get_counters()	167
7.29.3.6 operator()()	167
7.29.3.7 operator[]()	167
7.29.3.8 push_back()	167
7.29.3.9 reserve()	168
7.29.3.10 shrink_to_fit()	168
7.29.3.11 size()	168
7.30 PhyloRuleDynData Class Reference	168
7.30.1 Detailed Description	168
7.30.2 Constructor & Destructor Documentation	169
7.30.2.1 PhyloRuleDynData()	169
$7.30.2.2 \sim$ PhyloRuleDynData()	169
7.30.3 Member Data Documentation	169
7.30.3.1 counts	169
7.30.3.2 duplication	169
7.30.3.3 lb	169
7.30.3.4 pos	170

7.30.3.5 ub
7.31 PowerSet< Array_Type, Data_Rule_Type > Class Template Reference
7.31.1 Detailed Description
7.31.2 Constructor & Destructor Documentation
7.31.2.1 PowerSet() [1/3]
7.31.2.2 PowerSet() [2/3]
7.31.2.3 PowerSet() [3/3]
7.31.2.4 ~PowerSet()
7.31.3 Member Function Documentation
7.31.3.1 add_rule() [1/2]
7.31.3.2 add_rule() [2/2]
7.31.3.3 begin()
7.31.3.4 calc()
7.31.3.5 end()
7.31.3.6 get_data()
7.31.3.7 get_data_ptr()
7.31.3.8 init_support()
7.31.3.9 operator[]()
7.31.3.10 reset()
7.31.3.11 size()
7.31.4 Member Data Documentation
7.31.4.1 coordinates_free
7.31.4.2 coordinates_locked
7.31.4.3 data
7.31.4.4 EmptyArray
7.31.4.5 M
7.31.4.6 N
7.31.4.7 n_free
7.31.4.8 n_locked
7.31.4.9 rules
7.31.4.10 rules_deleted
7.32 Progress Class Reference
7.32.1 Detailed Description
7.32.2 Constructor & Destructor Documentation
7.32.2.1 Progress()
7.32.2.2 ~ Progress()
7.32.3 Member Function Documentation
7.32.3.1 end()
7.32.3.2 next()
7.33 Rule < Array_Type, Data_Type > Class Template Reference
7.33.1 Detailed Description
7.33.2 Constructor & Destructor Documentation

7.33.2.1 Rule() [1/2]	179
7.33.2.2 Rule() [2/2]	179
7.33.2.3 ~Rule()	180
7.33.3 Member Function Documentation	180
7.33.3.1 D()	180
7.33.3.2 operator()()	180
7.34 Rules< Array_Type, Data_Type > Class Template Reference	180
7.34.1 Detailed Description	181
7.34.2 Constructor & Destructor Documentation	181
7.34.2.1 Rules() [1/2]	181
7.34.2.2 Rules() [2/2]	181
7.34.2.3 ∼Rules()	182
7.34.3 Member Function Documentation	182
7.34.3.1 add_rule() [1/2]	182
7.34.3.2 add_rule() [2/2]	182
7.34.3.3 get_seq()	182
7.34.3.4 operator()()	183
7.34.3.5 operator=()	183
7.34.3.6 size()	183
7.35 StatsCounter< Array_Type, Data_Type > Class Template Reference	184
7.35.1 Detailed Description	184
7.35.2 Constructor & Destructor Documentation	184
7.35.2.1 StatsCounter() [1/3]	184
7.35.2.2 StatsCounter() [2/3]	185
7.35.2.3 StatsCounter() [3/3]	185
7.35.2.4 \sim StatsCounter()	185
7.35.3 Member Function Documentation	185
7.35.3.1 add_counter() [1/2]	185
7.35.3.2 add_counter() [2/2]	186
7.35.3.3 count_all()	186
7.35.3.4 count_current()	186
7.35.3.5 count_init()	186
7.35.3.6 get_counters()	186
7.35.3.7 get_descriptions()	186
7.35.3.8 get_names()	187
7.35.3.9 reset_array()	187
7.35.3.10 set_counters()	187
7.35.3.11 size()	187
7.36 Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > Class Tem-	
•	187
7.36.1 Detailed Description	
7.36.2 Constructor & Destructor Documentation	190

7.36.2.1 Support() [1/3]	190
7.36.2.2 Support() [2/3]	190
7.36.2.3 Support() [3/3]	190
7.36.2.4 ~Support()	190
7.36.3 Member Function Documentation	191
7.36.3.1 add_counter() [1/2]	191
7.36.3.2 add_counter() [2/2]	191
7.36.3.3 add_rule() [1/2]	191
7.36.3.4 add_rule() [2/2]	191
7.36.3.5 add_rule_dyn() [1/2]	191
7.36.3.6 add_rule_dyn() [2/2]	192
7.36.3.7 calc()	192
7.36.3.8 eval_rules_dyn()	192
7.36.3.9 get_counters()	192
7.36.3.10 get_counts()	193
7.36.3.11 get_current_stats()	193
7.36.3.12 get_data()	193
7.36.3.13 get_rules()	193
7.36.3.14 get_rules_dyn()	193
7.36.3.15 init_support()	194
7.36.3.16 print()	194
7.36.3.17 reset_array() [1/2]	194
7.36.3.18 reset_array() [2/2]	194
7.36.3.19 set_counters()	194
7.36.3.20 set_rules()	195
7.36.3.21 set_rules_dyn()	195
7.36.4 Member Data Documentation	195
7.36.4.1 change_stats	195
7.36.4.2 coordiantes_n_free	195
7.36.4.3 coordiantes_n_locked	195
7.36.4.4 coordinates_free	196
7.36.4.5 coordinates_locked	196
7.36.4.6 current_stats	196
7.36.4.7 delete_counters	196
7.36.4.8 delete_rules	196
7.36.4.9 delete_rules_dyn	197
7.36.4.10 hashes	197
7.36.4.11 hashes_initialized	197
7.36.4.12 M	197
	197
7.36.4.14 N	198
7.36.4.15 n. counters	198

	7.37 vecHasher $<$ T $>$ Struct Template Reference	198
	7.37.1 Detailed Description	198
	7.37.2 Member Function Documentation	198
	7.37.2.1 operator()()	198
ΩΙ	File Documentation	199
0 1	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 BARRAY_TEMPLATE	
	8.3.1.2 BARRAY TEMPLATE ARGS	
	8.3.1.3 BARRAY_TYPE	
	8.3.1.4 BARRY_BARRAY_MEAT_OPERATORS_HPP	
	8.3.1.5 COL	
	8.3.1.6 ROW	
	8.3.2 Function Documentation	
	8.3.2.1 BARRAY_TEMPLATE() [1/6]	
	8.3.2.2 BARRAY_TEMPLATE() [2/6]	
	8.3.2.4 BARRAY_TEMPLATE() [4/6]	
	8.3.2.5 BARRAY_TEMPLATE() [5/6]	
	8.3.2.6 BARRAY TEMPLATE() [6/6]	
	8.3.2.7 BARRAY TEMPLATE () [6/6]	
	8.3.2.8 BARRAY_TYPE()	
	_ ·	
	8.3.2.9 for()	
	8.3.2.10 operator()()	
	8.3.3.1 rhs	
	8.3.3.2 this	
	8.4 include/barry/barray-meat.hpp File Reference	
	8.4.1 Macro Definition Documentation	
	8.4.1.1 BARRAY TEMPLATE	
	8.4.1.2 BARRAY TEMPLATE ARGS	
	8.4.1.3 BARRAY_TYPE	
	8.4.1.4 COL	
	8.4.1.5 ROW	
	8.4.2 Function Documentation	
	8.4.2.1 ans()	
	U.T.C.C UMINAL ILIVII LALLU 11/7.01	∠ U.9

8.4.2.3 BARRAY_TEMPLATE() [2/23]
8.4.2.4 BARRAY_TEMPLATE() [3/23]
8.4.2.5 BARRAY_TEMPLATE() [4/23]
8.4.2.6 BARRAY_TEMPLATE() [5/23]
8.4.2.7 BARRAY_TEMPLATE() [6/23]
8.4.2.8 BARRAY_TEMPLATE() [7/23]
8.4.2.9 BARRAY_TEMPLATE() [8/23]
8.4.2.10 BARRAY_TEMPLATE() [9/23]
8.4.2.11 BARRAY_TEMPLATE() [10/23]
8.4.2.12 BARRAY_TEMPLATE() [11/23]
8.4.2.13 BARRAY_TEMPLATE() [12/23]
8.4.2.14 BARRAY_TEMPLATE() [13/23]
8.4.2.15 BARRAY_TEMPLATE() [14/23]
8.4.2.16 BARRAY_TEMPLATE() [15/23]
8.4.2.17 BARRAY_TEMPLATE() [16/23]
8.4.2.18 BARRAY_TEMPLATE() [17/23]
8.4.2.19 BARRAY_TEMPLATE() [18/23]
8.4.2.20 BARRAY_TEMPLATE() [19/23]
8.4.2.21 BARRAY_TEMPLATE() [20/23]
8.4.2.22 BARRAY_TEMPLATE() [21/23]
8.4.2.23 BARRAY_TEMPLATE() [22/23]
8.4.2.24 BARRAY_TEMPLATE() [23/23]
8.4.2.25 COL()
8.4.2.26 for() [1/3]
8.4.2.27 for() [2/3]
8.4.2.28 for() [3/3]
8.4.2.29 if() [1/17]
8.4.2.30 if() [2/17]
8.4.2.31 if() [3/17]
8.4.2.32 if() [4/17]
8.4.2.33 if() [5/17]
8.4.2.34 if() [6/17]
8.4.2.35 if() [7/17]
8.4.2.36 if() [8/17]
8.4.2.37 if() [9/17]
8.4.2.38 if() [10/17]
8.4.2.39 if() [11/17]
8.4.2.40 if() [12/17]
8.4.2.41 if() [13/17]
8.4.2.42 if() [14/17]
8.4.2.43 if() [15/17]
8.4.2.44 if() [16/17]

8.4.2.45 if() [17/17]	217
8.4.2.46 M()	217
8.4.2.47 resize() [1/2]	217
8.4.2.48 resize() [2/2]	217
8.4.2.49 return()	217
8.4.2.50 ROW() [1/2]	217
8.4.2.51 ROW() [2/2]	217
8.4.3 Variable Documentation	218
8.4.3.1 add	218
8.4.3.2 ans	218
8.4.3.3 Array	218
8.4.3.4 check_bounds	218
8.4.3.5 check_exists	219
8.4.3.6 col0	219
8.4.3.7 const	219
8.4.3.8 copy_data	219
8.4.3.9 data	219
8.4.3.10 delete_data	220
8.4.3.11 delete_data	220
8.4.3.12 else	220
8.4.3.13 false	220
8.4.3.14 first	220
8.4.3.15 i1	221
8.4.3.16 j	221
8.4.3.17 j0	221
8.4.3.18 j1	221
8.4.3.19 M	221
8.4.3.20 M	222
8.4.3.21 N	222
8.4.3.22 NCells	222
8.4.3.23 report	222
8.4.3.24 return	222
8.4.3.25 row0	223
8.4.3.26 search	223
8.4.3.27 source	223
8.4.3.28 target	223
8.4.3.29 v	223
8.4.3.30 value	223
8.5 include/barry/barraycell-bones.hpp File Reference	224
8.6 include/barry/barraycell-meat.hpp File Reference	224
8.7 include/barry/barraydense-bones.hpp File Reference	225
8.7.1 Macro Definition Documentation	227

8.7.1.1 BARRY_BARRAYDENSE_BONES_HPP	227
8.8 include/barry/barraydense-meat-operators.hpp File Reference	227
8.8.1 Macro Definition Documentation	228
8.8.1.1 BARRY_BARRAYDENSE_MEAT_OPERATORS_HPP	228
8.8.1.2 BDENSE_TEMPLATE	228
8.8.1.3 BDENSE_TEMPLATE_ARGS	228
8.8.1.4 BDENSE_TYPE	229
8.8.1.5 COL	229
8.8.1.6 POS	229
8.8.1.7 POS_N	229
8.8.1.8 ROW	229
8.8.2 Function Documentation	229
8.8.2.1 BDENSE_TEMPLATE() [1/4]	230
8.8.2.2 BDENSE_TEMPLATE() [2/4]	230
8.8.2.3 BDENSE_TEMPLATE() [3/4]	230
8.8.2.4 BDENSE_TEMPLATE() [4/4]	230
8.8.2.5 BDENSE_TEMPLATE_ARGS()	230
8.8.2.6 BDENSE_TYPE()	230
8.9 include/barry/barraydense-meat.hpp File Reference	231
8.9.1 Macro Definition Documentation	233
8.9.1.1 BDENSE_TEMPLATE	233
8.9.1.2 BDENSE_TEMPLATE_ARGS	234
8.9.1.3 BDENSE_TYPE	
8.9.1.4 COL	234
8.9.1.5 POS	234
8.9.1.6 POS_N	234
8.9.1.7 ROW	235
8.9.1.8 ZERO_CELL	235
8.9.2 Function Documentation	
8.9.2.1 ans()	235
8.9.2.2 BDENSE_TEMPLATE() [1/37]	
8.9.2.3 BDENSE_TEMPLATE() [2/37]	235
8.9.2.4 BDENSE_TEMPLATE() [3/37]	235
8.9.2.5 BDENSE_TEMPLATE() [4/37]	
8.9.2.6 BDENSE_TEMPLATE() [5/37]	236
8.9.2.7 BDENSE_TEMPLATE() [6/37]	236
8.9.2.8 BDENSE_TEMPLATE() [7/37]	236
8.9.2.9 BDENSE_TEMPLATE() [8/37]	
8.9.2.10 BDENSE_TEMPLATE() [9/37]	
8.9.2.11 BDENSE_TEMPLATE() [10/37]	
8.9.2.12 BDENSE_TEMPLATE() [11/37]	
8.9.2.13 BDENSE_TEMPLATE() [12/37]	237

8.9.2.14 BDENSE_TEMPLATE() [13/37]
8.9.2.15 BDENSE_TEMPLATE() [14/37]
8.9.2.16 BDENSE_TEMPLATE() [15/37]
8.9.2.17 BDENSE_TEMPLATE() [16/37]
8.9.2.18 BDENSE_TEMPLATE() [17/37]
8.9.2.19 BDENSE_TEMPLATE() [18/37]
8.9.2.20 BDENSE_TEMPLATE() [19/37]
8.9.2.21 BDENSE_TEMPLATE() [20/37]
8.9.2.22 BDENSE_TEMPLATE() [21/37]
8.9.2.23 BDENSE_TEMPLATE() [22/37]
8.9.2.24 BDENSE_TEMPLATE() [23/37]
8.9.2.25 BDENSE_TEMPLATE() [24/37]
8.9.2.26 BDENSE_TEMPLATE() [25/37]
8.9.2.27 BDENSE_TEMPLATE() [26/37]
8.9.2.28 BDENSE_TEMPLATE() [27/37]
8.9.2.29 BDENSE_TEMPLATE() [28/37]
8.9.2.30 BDENSE_TEMPLATE() [29/37]
8.9.2.31 BDENSE_TEMPLATE() [30/37]
8.9.2.32 BDENSE_TEMPLATE() [31/37]
8.9.2.33 BDENSE_TEMPLATE() [32/37]
8.9.2.34 BDENSE_TEMPLATE() [33/37]
8.9.2.35 BDENSE_TEMPLATE() [34/37]
8.9.2.36 BDENSE_TEMPLATE() [35/37]
8.9.2.37 BDENSE_TEMPLATE() [36/37]
8.9.2.38 BDENSE_TEMPLATE() [37/37]
8.9.2.39 for()
8.9.2.40 if() [1/4]
8.9.2.41 if() [2/4]
8.9.2.42 if() [3/4]
8.9.2.43 if() [4/4]
8.9.2.44 insert_cell() [1/2]
8.9.2.45 insert_cell() [2/2]
8.9.2.46 M()
8.9.2.47 resize() [1/6]
8.9.2.48 resize() [2/6]
8.9.2.49 resize() [3/6]
8.9.2.50 resize() [4/6]
8.9.2.51 resize() [5/6]
8.9.2.52 resize() [6/6]
8.9.2.53 rm_cell() [1/3]
8.9.2.54 rm_cell() [2/3]
24 2 22 cm (COM) 13/31 24

8.9.2.56 va_end()	 245
8.9.2.57 va_start()	 245
8.9.2.58 vprintf()	 245
8.9.3 Variable Documentation	 245
8.9.3.1 add	 246
8.9.3.2 ans	 246
8.9.3.3 check_bounds	 246
8.9.3.4 check_exists	 246
8.9.3.5 col	 247
8.9.3.6 const	 247
8.9.3.7 copy_data	 247
8.9.3.8 data	 247
8.9.3.9 delete_data	 247
8.9.3.10 delete_data	 248
8.9.3.11 el	 248
8.9.3.12 el_colsums	 248
8.9.3.13 el_rowsums	 248
8.9.3.14 else	 248
8.9.3.15 false	 249
8.9.3.16 i1	 249
8.9.3.17 j	 249
8.9.3.18 j0	 249
8.9.3.19 j1	 249
8.9.3.20 M	 249
8.9.3.21 M	 250
8.9.3.22 N	
8.9.3.23 report	 250
8.9.3.24 return	 250
8.9.3.25 source	 250
8.9.3.26 target	 251
8.9.3.27 v	 251
8.9.3.28 val0	 251
8.9.3.29 val1	 251
8.9.3.30 value	 251
8.10 include/barry/barraydensecell-bones.hpp File Reference	
8.10.1 Macro Definition Documentation	 252
8.10.1.1 POS	 253
8.11 include/barry/barraydensecell-meat.hpp File Reference	 253
8.11.1 Macro Definition Documentation	
8.11.1.1 POS	
8.12 include/barry/barraydensecol-bones.hpp File Reference	 254
8.12.1 Macro Definition Documentation	255

8.12.1.1 POS	55
8.12.1.2 POS_N	55
8.12.1.3 ZERO_CELL	55
8.13 include/barry/barraydenserow-bones.hpp File Reference	55
8.13.1 Macro Definition Documentation	56
8.13.1.1 POS	56
8.13.1.2 POS_N	56
8.13.1.3 ZERO_CELL	56
8.14 include/barry/barrayrow-bones.hpp File Reference	57
8.15 include/barry/barrayrow-meat.hpp File Reference	57
8.15.1 Macro Definition Documentation	58
8.15.1.1 BARRY_BARRAYROW_MEAT_HPP	58
8.15.1.2 BROW_TEMPLATE	59
8.15.1.3 BROW_TEMPLATE_ARGS	59
8.15.1.4 BROW_TYPE	59
8.15.2 Function Documentation	59
8.15.2.1 BROW_TEMPLATE() [1/5]	59
8.15.2.2 BROW_TEMPLATE() [2/5]	59
8.15.2.3 BROW_TEMPLATE() [3/5]	30
8.15.2.4 BROW_TEMPLATE() [4/5]	30
8.15.2.5 BROW_TEMPLATE() [5/5]	30
8.16 include/barry/barrayvector-bones.hpp File Reference	30
8.17 include/barry/barrayvector-meat.hpp File Reference	31
8.17.1 Macro Definition Documentation	32
8.17.1.1 BARRY_BARRAYVECTOR_MEAT_HPP	32
8.18 include/barry/barry-configuration.hpp File Reference	32
8.18.1 Macro Definition Documentation	33
8.18.1.1 BARRY_CHECK_SUPPORT	33
8.18.1.2 BARRY_ISFINITE	33
8.18.1.3 BARRY_MAX_NUM_ELEMENTS	33
8.18.1.4 BARRY_SAFE_EXP	33
8.18.1.5 printf_barry	33
8.18.2 Typedef Documentation	33
8.18.2.1 Map	34
8.19 include/barry/barry-debug.hpp File Reference	34
8.19.1 Macro Definition Documentation	34
8.19.1.1 BARRY_DEBUG_LEVEL	34
8.20 include/barry/barry-macros.hpp File Reference	34
8.20.1 Macro Definition Documentation	35
8.20.1.1 BARRY_ONE	35
8.20.1.2 BARRY_ONE_DENSE	35
8.20.1.3 BARRY UNUSED	35

8.20.1.4 BARRY_ZERO	35
8.20.1.5 BARRY_ZERO_DENSE	36
8.21 include/barry/barry.hpp File Reference	36
8.21.1 Macro Definition Documentation	37
8.21.1.1 BARRY_HPP	37
8.21.1.2 BARRY_VERSION	37
8.21.1.3 COUNTER_FUNCTION	38
8.21.1.4 COUNTER_LAMBDA	38
8.21.1.5 RULE_FUNCTION	38
8.21.1.6 RULE_LAMBDA	38
8.22 include/barry/cell-bones.hpp File Reference	39
8.23 include/barry/cell-meat.hpp File Reference	39
8.24 include/barry/col-bones.hpp File Reference	70
8.25 include/barry/counters-bones.hpp File Reference	70
8.26 include/barry/counters-meat.hpp File Reference	72
8.26.1 Macro Definition Documentation	73
8.26.1.1 COUNTER_TEMPLATE	74
8.26.1.2 COUNTER_TEMPLATE_ARGS	74
8.26.1.3 COUNTER_TYPE	74
8.26.1.4 COUNTERS_TEMPLATE	74
8.26.1.5 COUNTERS_TEMPLATE_ARGS	74
8.26.1.6 COUNTERS_TYPE	74
8.26.2 Function Documentation	75
8.26.2.1 count_fun()	75
8.26.2.2 COUNTER_TEMPLATE() [1/7]	75
8.26.2.3 COUNTER_TEMPLATE() [2/7]	75
8.26.2.4 COUNTER_TEMPLATE() [3/7]	75
8.26.2.5 COUNTER_TEMPLATE() [4/7]	75
8.26.2.6 COUNTER_TEMPLATE() [5/7]	76
8.26.2.7 COUNTER_TEMPLATE() [6/7]	76
8.26.2.8 COUNTER_TEMPLATE() [7/7]	76
8.26.2.9 Counters()	76
8.26.2.10 COUNTERS_TEMPLATE() [1/7]	76
8.26.2.11 COUNTERS_TEMPLATE() [2/7]	77
8.26.2.12 COUNTERS_TEMPLATE() [3/7]	77
8.26.2.13 COUNTERS_TEMPLATE() [4/7]	77
8.26.2.14 COUNTERS_TEMPLATE() [5/7]	77
8.26.2.15 COUNTERS_TEMPLATE() [6/7]	77
8.26.2.16 COUNTERS_TEMPLATE() [7/7]	78
8.26.2.17 data()	78
8.26.2.18 desc()	78
8.26.2.19 init_fun() [1/3]	78

8.26.2.20 init_fun() [2/3]	:78
8.26.2.21 init_fun() [3/3]	78
8.26.2.22 name()	:79
8.26.3 Variable Documentation	:79
8.26.3.1 count_fun	:79
8.26.3.2 counter	79
8.26.3.3 counter	:79
8.26.3.4 data	:80
8.26.3.5 desc	:80
8.26.3.6 i	:80
8.26.3.7 init_fun	:80
8.26.3.8 j	:80
8.26.3.9 name	81
8.26.3.10 noexcept	81
8.26.3.11 return	81
8.26.3.12 this	:81
8.27 include/barry/counters/network-css.hpp File Reference	82
8.27.1 Macro Definition Documentation	:83
8.27.1.1 CSS_APPEND	:83
8.27.1.2 CSS_CASE_ELSE	84
8.27.1.3 CSS_CASE_PERCEIVED	84
8.27.1.4 CSS_CASE_TRUTH	84
8.27.1.5 CSS_CHECK_SIZE	84
8.27.1.6 CSS_CHECK_SIZE_INIT	84
8.27.1.7 CSS_NET_COUNTER_LAMBDA_INIT	:85
8.27.1.8 CSS_PERCEIVED_CELLS	85
8.27.1.9 CSS_SIZE	:85
8.27.1.10 CSS_TRUE_CELLS	85
8.27.2 Function Documentation	:85
8.27.2.1 counter_css_census01()	:86
8.27.2.2 counter_css_census02()	:86
8.27.2.3 counter_css_census03()	:86
8.27.2.4 counter_css_census04()	:86
8.27.2.5 counter_css_census05()	87
8.27.2.6 counter_css_census06()	:87
8.27.2.7 counter_css_census07()	:87
8.27.2.8 counter_css_census08()	87
8.27.2.9 counter_css_census09()	:88
8.27.2.10 counter_css_census10()	:88
8.27.2.11 counter_css_completely_false_recip_comiss()	88
8.27.2.12 counter_css_completely_false_recip_omiss()	:88
8.27.2.13 counter_css_mixed_recip()	89

8.27.2.14 counter_css_partially_false_recip_commi()	289
8.27.2.15 counter_css_partially_false_recip_omiss()	289
8.28 include/barry/counters/network.hpp File Reference	290
8.28.1 Macro Definition Documentation	292
8.28.1.1 BARRY_ZERO_NETWORK	293
8.28.1.2 BARRY_ZERO_NETWORK_DENSE	293
8.28.1.3 NET_C_DATA_IDX	293
8.28.1.4 NET_C_DATA_NUM	293
8.28.1.5 NETWORK_COUNTER	293
8.28.1.6 NETWORK_COUNTER_LAMBDA	294
8.28.1.7 NETWORK_RULE	294
8.28.1.8 NETWORK_RULE_LAMBDA	294
8.28.1.9 NETWORKDENSE_COUNTER_LAMBDA	294
8.28.2 Typedef Documentation	295
8.28.2.1 NetCounter	295
8.28.2.2 NetCounters	295
8.28.2.3 NetModel	295
8.28.2.4 NetRule	295
8.28.2.5 NetRules	295
8.28.2.6 NetStatsCounter	296
8.28.2.7 NetSupport	296
8.28.2.8 Network	296
8.28.2.9 NetworkDense	296
8.28.3 Function Documentation	296
8.28.3.1 rules_zerodiag()	296
8.29 include/barry/counters/phylo.hpp File Reference	297
8.29.1 Macro Definition Documentation	299
8.29.1.1 DEFAULT_DUPLICATION	299
8.29.1.2 DUPL_DUPL	299
8.29.1.3 DUPL_EITH	299
8.29.1.4 DUPL_SPEC	300
8.29.1.5 IF_MATCHES	300
8.29.1.6 IF_NOTMATCHES	300
8.29.1.7 IS_DUPLICATION	300
8.29.1.8 IS_EITHER	300
8.29.1.9 IS_SPECIATION	301
8.29.1.10 MAKE_DUPL_VARS	301
8.29.1.11 PHYLO_CHECK_MISSING	301
8.29.1.12 PHYLO_COUNTER_LAMBDA	301
8.29.1.13 PHYLO_RULE_DYN_LAMBDA	302
8.29.2 Typedef Documentation	302
8.29.2.1 PhyloArray	302

8.29.2.2 PhyloCounter
8.29.2.3 PhyloCounters
8.29.2.4 PhyloModel
8.29.2.5 PhyloPowerSet
8.29.2.6 PhyloRule
8.29.2.7 PhyloRuleData
8.29.2.8 PhyloRuleDyn
8.29.2.9 PhyloRules
8.29.2.10 PhyloRulesDyn
8.29.2.11 PhyloStatsCounter
8.29.2.12 PhyloSupport
8.29.3 Function Documentation
8.29.3.1 get_last_name()
8.30 include/barry/model-bones.hpp File Reference
8.30.1 Function Documentation
8.30.1.1 keygen_default()
8.31 include/barry/model-meat.hpp File Reference
8.31.1 Macro Definition Documentation
8.31.1.1 MODEL_TEMPLATE
8.31.1.2 MODEL_TEMPLATE_ARGS
8.31.1.3 MODEL_TYPE
8.31.2 Function Documentation
8.31.2.1 likelihood_()
8.31.2.2 MODEL_TEMPLATE() [1/2]
8.31.2.3 MODEL_TEMPLATE() [2/2]
8.31.2.4 update_normalizing_constant()
8.32 include/barry/models/defm.hpp File Reference
8.33 include/barry/models/defm/defm-bones.hpp File Reference
8.33.1 Macro Definition Documentation
8.33.1.1 INITIALIZED
8.33.2 Function Documentation
8.33.2.1 keygen_full()
8.33.2.2 RULE_FUNCTION()
8.33.2.3 vec_diff()
8.34 include/barry/models/defm/defm-meat.hpp File Reference
8.35 include/barry/models/geese.hpp File Reference
8.36 include/barry/models/geese/flock-bones.hpp File Reference
8.37 include/barry/models/geese/flock-meat.hpp File Reference
8.38 include/barry/models/geese/geese-bones.hpp File Reference
8.38.1 Macro Definition Documentation
8.38.1.1 INITIALIZED
8.38.2 Function Documentation

8.38.2.1 keygen_full()	13
8.38.2.2 RULE_FUNCTION()	13
8.38.2.3 vec_diff()	14
8.38.2.4 vector_caster()	14
8.39 include/barry/models/geese/geese-meat-constructors.hpp File Reference	14
8.40 include/barry/models/geese/geese-meat-likelihood.hpp File Reference	15
8.41 include/barry/models/geese/geese-meat-likelihood_exhaust.hpp File Reference	16
8.42 include/barry/models/geese/geese-meat-predict.hpp File Reference	16
8.43 include/barry/models/geese/geese-meat-predict_exhaust.hpp File Reference	17
8.44 include/barry/models/geese/geese-meat-predict_sim.hpp File Reference	17
8.45 include/barry/models/geese/geese-meat-simulate.hpp File Reference	18
8.46 include/barry/models/geese/geese-meat.hpp File Reference	18
8.47 include/barry/models/geese/geese-node-bones.hpp File Reference	19
8.48 include/barry/powerset-bones.hpp File Reference	19
8.49 include/barry/powerset-meat.hpp File Reference	20
8.50 include/barry/progress.hpp File Reference	22
8.50.1 Macro Definition Documentation	22
8.50.1.1 BARRY_PROGRESS_BAR_WIDTH	22
8.51 include/barry/rules-bones.hpp File Reference	22
8.51.1 Function Documentation	23
8.51.1.1 rule_fun_default()	23
8.52 include/barry/rules-meat.hpp File Reference	24
8.53 include/barry/statscounter-bones.hpp File Reference	24
8.54 include/barry/statscounter-meat.hpp File Reference	26
8.54.1 Macro Definition Documentation	27
8.54.1.1 STATSCOUNTER_TEMPLATE	27
8.54.1.2 STATSCOUNTER_TEMPLATE_ARGS	27
8.54.1.3 STATSCOUNTER_TYPE	28
8.54.2 Function Documentation	28
8.54.2.1 clear()	28
8.54.2.2 for()	28
8.54.2.3 resize()	28
8.54.2.4 STATSCOUNTER_TEMPLATE() [1/9]	28
8.54.2.5 STATSCOUNTER_TEMPLATE() [2/9]	28
8.54.2.6 STATSCOUNTER_TEMPLATE() [3/9]	29
8.54.2.7 STATSCOUNTER_TEMPLATE() [4/9]	29
8.54.2.8 STATSCOUNTER_TEMPLATE() [5/9]	29
8.54.2.9 STATSCOUNTER_TEMPLATE() [6/9]	29
8.54.2.10 STATSCOUNTER_TEMPLATE() [7/9]	29
8.54.2.11 STATSCOUNTER_TEMPLATE() [8/9]	29
8.54.2.12 STATSCOUNTER_TEMPLATE() [9/9]	30
8.54.3 Variable Documentation 33	30

8.54.3.1 counter	30
8.54.3.2 counter_deleted	30
8.54.3.3 counters	30
8.54.3.4 counters	30
8.54.3.5 current_stats	31
8.54.3.6 EmptyArray	31
8.54.3.7 f	31
8.54.3.8 j	31
8.54.3.9 return	31
8.55 include/barry/statsdb.hpp File Reference	32
8.56 include/barry/support-bones.hpp File Reference	32
8.57 include/barry/support-meat.hpp File Reference	34
8.57.1 Macro Definition Documentation	35
8.57.1.1 BARRY_SUPPORT_MEAT_HPP	36
8.57.1.2 SUPPORT_TEMPLATE	36
8.57.1.3 SUPPORT_TEMPLATE_ARGS	36
8.57.1.4 SUPPORT_TYPE	36
8.57.2 Function Documentation	36
8.57.2.1 calc_backend_dense()	37
8.57.2.2 calc_backend_sparse()	37
8.57.2.3 for()	37
8.57.2.4 if() [1/4]	37
8.57.2.5 if() [2/4]	37
8.57.2.6 if() [3/4]	37
8.57.2.7 if() [4/4]	38
8.57.2.8 insert_cell() [1/2]	38
8.57.2.9 insert_cell() [2/2]	38
8.57.2.10 rm_cell()	38
8.57.2.11 SUPPORT_TEMPLATE() [1/17]	38
8.57.2.12 SUPPORT_TEMPLATE() [2/17]	39
8.57.2.13 SUPPORT_TEMPLATE() [3/17]	39
8.57.2.14 SUPPORT_TEMPLATE() [4/17]	39
8.57.2.15 SUPPORT_TEMPLATE() [5/17]	39
8.57.2.16 SUPPORT_TEMPLATE() [6/17]	39
8.57.2.17 SUPPORT_TEMPLATE() [7/17]	10
8.57.2.18 SUPPORT_TEMPLATE() [8/17]	10
8.57.2.19 SUPPORT_TEMPLATE() [9/17]	10
8.57.2.20 SUPPORT_TEMPLATE() [10/17]	1 0
8.57.2.21 SUPPORT_TEMPLATE() [11/17]	1 0
8.57.2.22 SUPPORT_TEMPLATE() [12/17]	1 0
8.57.2.23 SUPPORT_TEMPLATE() [13/17]	11
8.57.2.24 SUPPORT_TEMPLATE() [14/17]	11

8.57.2.25 SUPPORT_TEMPLATE() [15/17]	341
8.57.2.26 SUPPORT_TEMPLATE() [16/17]	341
8.57.2.27 SUPPORT_TEMPLATE() [17/17]	341
8.57.3 Variable Documentation	341
8.57.3.1 array_bank	342
8.57.3.2 change_stats_different	342
8.57.3.3 coord_i	342
8.57.3.4 coord_j	342
8.57.3.5 counters	342
8.57.3.6 counters	342
8.57.3.7 delete_counters	343
8.57.3.8 delete_rules	343
8.57.3.9 delete_rules_dyn	343
8.57.3.10 else	343
8.57.3.11 f	343
8.57.3.12 hashes	344
8.57.3.13 return	344
8.57.3.14 rules	344
8.57.3.15 rules	344
8.57.3.16 rules_dyn	344
8.57.3.17 stats_bank	345
8.57.3.18 tmp_chng	345
8.58 include/barry/typedefs.hpp File Reference	345
8.58.1 Typedef Documentation	347
8.58.1.1 Col_type	347
8.58.1.2 Counter_fun_type	347
8.58.1.3 Counts_type	348
8.58.1.4 MapVec_type	348
8.58.1.5 Row_type	348
8.58.1.6 Rule_fun_type	348
8.58.1.7 uint	348
8.58.2 Function Documentation	348
8.58.2.1 vec_equal()	348
8.58.2.2 vec_equal_approx()	349
8.58.2.3 vec_inner_prod() [1/2]	349
8.58.2.4 vec_inner_prod() [2/2]	349
8.59 README.md File Reference	349
Index	351

Main Page

Barry: your to-go motif accountant

This repository contains a C++ template library that essentially counts sufficient statistics on binary arrays. The goal of the library is to provide a general framework for building discrete exponential-family models. A particular example is Exponential Random Graph Models (ERGMs), but we can use barry to deal with non-square arrays.

Among the key features included in barry, we have:

- · Sparse arrays.
- · User-defined count statistics.
- · User-defined constrain of the support set.
- · Powerset generation of binary arrays.
- Discrete Exponential Family Models module (DEFMs).
- · Pooled DEFMs.

This was created and maintained by Dr. George G. Vega Yon as part of his doctoral dissertation "Essays on Bioinformatics and Social Network Analysis: Statistical and Computational Methods for Complex Systems."

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() {
```

2 Main Page

```
// 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?
  net.print("Current view");
  // Adding extra ties
  net += {1, 0};
net(2, 0) = true;
  // And removing a couple
  net(0, 0) = false;
net -= {1, 1};
  net.print("New view");
  // 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);
  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 «
                         : " « counts[0] « std::endl «
    "Transitive triads : " « counts[U] « std::endl «
"Transitive triads : " « counts[1] « std::endl «
"Isolates : " « counts[2] « std::endl «
                         : " « counts[3] « std::endl « : " « counts[4] « std::endl;
    "C triads
    "Mutuals
  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
               1
   1,]
            1
   2,] .
                   . 1
   3,] . . . . . . 4,] 1 . 1 .
   5,]
  0,] .
1,] 1
           1
               1
  2,] 1 .
                      1
   3,] . . .
4,] 1 . 1
   5,] . . . .
Edges
Transitive triads : 3
Isolates
C triads
Mutuals
```

Features

Efficient memory usage

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

Documentation

More information can be found in the Doxygen website here and in the PDF version of the documentation here.

Code of Conduct

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

4 Main Page

Module Index

2.1 Modules

Here is a list of all modules:

Counting					 						 	 											1
Statistical Models											 	 											1
Network counters					 						 	 											12
Phylo counters					 						 	 											19
Phylo rules																							27

6 Module Index

Class Index

3.1 Class List

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

BArray< Cell_Type, Data_Type >
Baseline class for binary arrays
BArrayCell < Cell_Type, Data_Type >
BArrayCell_const< Cell_Type, Data_Type >
BArrayDense < Cell_Type, Data_Type >
Baseline class for binary arrays
BArrayDenseCell< Cell_Type, Data_Type >
BArrayDenseCell_const< Cell_Type, Data_Type >
BArrayDenseCol< Cell_Type, Data_Type >
BArrayDenseCol_const< Cell_Type, Data_Type >
BArrayDenseRow< Cell_Type, Data_Type >
BArrayDenseRow_const< Cell_Type, Data_Type >
BArrayRow< Cell_Type, Data_Type >
BArrayRow_const < Cell_Type, Data_Type >
BArrayVector< Cell_Type, Data_Type >
Row or column of a BArray
BArrayVector_const < Cell_Type, Data_Type >
Cell< Cell_Type >
Entries in BArray. For now, it only has two members:
Cell_const< Cell_Type >
ConstBArrayRowlter < Cell_Type, Data_Type >
Counter< Array_Type, Data_Type >
A counter function based on change statistics
Counters< Array_Type, Data_Type >
Vector of counters
Entries < Cell_Type >
A wrapper class to store source, target, val from a BArray object
Flock
A Flock is a group of Geese
FreqTable < T >
Frequency table of vectors
Geese
Annotated Phylo Model
Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >
General framework for discrete exponential models. This class allows generating discrete expo-
nential models in the form of a linear exponential model:

8 Class Index

NetCounterData	
Data class used to store arbitrary uint or double vectors	 154
NetworkData	
Data class for Networks	 156
Node	
A single node for the model	 158
NodeData	
Data definition for the PhyloArray class	 164
PhyloCounterData	 165
PhyloRuleDynData	
PowerSet< Array_Type, Data_Rule_Type >	
Powerset of a binary array	 170
Progress	
A simple progress bar	 177
Rule < Array_Type, Data_Type >	
Rule for determining if a cell should be included in a sequence	 178
Rules < Array_Type, Data_Type >	
Vector of objects of class Rule	 180
StatsCounter< Array_Type, Data_Type >	
Count stats for a single Array	 184
Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >	
Compute the support of sufficient statistics	 187
vecHasher <t></t>	 198

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-meat-operators.hpp
include/barry/barraydense-meat.hpp
include/barry/barraydensecell-bones.hpp
include/barry/barraydensecell-meat.hpp
include/barry/barraydensecol-bones.hpp
include/barry/barraydenserow-bones.hpp
include/barry/barrayrow-bones.hpp
include/barry/barrayrow-meat.hpp
include/barry/barrayvector-bones.hpp
include/barry/barrayvector-meat.hpp
include/barry/barry-configuration.hpp
include/barry/barry-debug.hpp
include/barry/barry-macros.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/progress.hpp
include/barry/rules-bones.hpp
include/barry/rules-meat.hpp
include/barry/statscounter-bones.hpp
include/barry/statscounter-meat.hpp

10 File Index

include/barry/statsdb.hpp
include/barry/support-bones.hpp
include/barry/support-meat.hpp 334
include/barry/typedefs.hpp
include/barry/counters/network-css.hpp
include/barry/counters/network.hpp
include/barry/counters/phylo.hpp
include/barry/models/defm.hpp
include/barry/models/geese.hpp
include/barry/models/defm/defm-bones.hpp
include/barry/models/defm/defm-meat.hpp
$include/barry/models/geese/flock-bones.hpp \\ \dots \\$
include/barry/models/geese/flock-meat.hpp
include/barry/models/geese/geese-bones.hpp
include/barry/models/geese/geese-meat-constructors.hpp
include/barry/models/geese/geese-meat-likelihood.hpp
include/barry/models/geese/geese-meat-likelihood_exhaust.hpp
include/barry/models/geese/geese-meat-predict.hpp
include/barry/models/geese/geese-meat-predict_exhaust.hpp
include/barry/models/geese/geese-meat-predict_sim.hpp
include/barry/models/geese/geese-meat-simulate.hpp
include/barry/models/geese/geese-meat.hpp
include/barry/models/geese/geese-node-bones.hpp

Module Documentation

5.1 Counting

Classes

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

Data definition for the PhyloArray class.

class Counter< Array_Type, Data_Type >

A counter function based on change statistics.

5.1.1 Detailed Description

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

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

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

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

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

5.2 Statistical Models

Statistical models available in barry.

Classes

class Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >

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

class Geese

Annotated Phylo Model.

class Flock

A Flock is a group of Geese.

5.2.1 Detailed Description

Statistical models available in barry.

5.3 Network counters

Counters for network models.

Functions

```
    template<typename Tnet = Network>
        void counter_edges (NetCounters< Tnet > *counters)
```

Number of edges.

template<typename Tnet = Network>

void counter_isolates (NetCounters< Tnet > *counters)

Number of isolated vertices.

- $\bullet \ \ template<> void \ counter_isolates \ (NetCounters< \ NetworkDense > *counters)$
- template<typename Tnet = Network>

```
void counter_mutual (NetCounters< Tnet > *counters)
```

Number of mutual ties.

- template<typename Tnet = Network>
 - void counter_istar2 (NetCounters< Tnet > *counters)
- template<> void counter_istar2 (NetCounters< NetworkDense > *counters)
- template<typename Tnet = Network>
 - void counter_ostar2 (NetCounters< Tnet > *counters)
- template<> void counter_ostar2 (NetCounters< NetworkDense > *counters)
- template<typename Tnet = Network>
 - void counter_ttriads (NetCounters< Tnet > *counters)
- template<> void counter_ttriads (NetCounters< NetworkDense > *counters)
- template<typename Tnet = Network>
 - void counter_ctriads (NetCounters< Tnet > *counters)
- template<> void counter_ctriads (NetCounters< NetworkDense > *counters)
- template<typename Tnet = Network>
 - void counter_density (NetCounters < Tnet > *counters)
- template<typename Tnet = Network>
 - void counter_idegree15 (NetCounters< Tnet > *counters)
- template<> void counter idegree15 (NetCounters< NetworkDense > *counters)
- template<typename Tnet = Network>
 - void counter_odegree15 (NetCounters< Tnet > *counters)
- template<> void counter_odegree15 (NetCounters< NetworkDense > *counters)

5.3 Network counters 13

```
• template<typename Tnet = Network>
  void counter_absdiff (NetCounters < Tnet > *counters, uint attr_id, double alpha=1.0)
     Sum of absolute attribute difference between ego and alter.
• template<typename Tnet = Network>
  void counter_diff (NetCounters < Tnet > *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)

• template<typename Tnet = Network>
  void counter nodeicov (NetCounters < Tnet > *counters, uint attr id)
• template<typename Tnet = Network>
  void counter nodeocov (NetCounters< Tnet > *counters, uint attr id)
template<typename Tnet = Network>
  void counter_nodecov (NetCounters< Tnet > *counters, uint attr_id)
• template<typename Tnet = Network>
  void counter_nodematch (NetCounters< Tnet > *counters, uint attr_id)
• template<typename Tnet = Network>
  void counter_idegree (NetCounters< Tnet > *counters, std::vector< uint > d)
     Counts number of vertices with a given in-degree.

    template<> void counter_idegree (NetCounters< NetworkDense > *counters, std::vector< uint > d)

• template<typename Tnet = Network>
  void counter odegree (NetCounters < Tnet > *counters, std::vector < uint > d)
     Counts number of vertices with a given out-degree.

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

template<typename Tnet = Network>
  void counter_degree (NetCounters< Tnet > *counters, std::vector< uint > d)
     Counts number of vertices with a given out-degree.
```

5.3.1 Detailed Description

Counters for network models.

Parameters

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

5.3.2 Function Documentation

5.3.2.1 counter_absdiff()

Sum of absolute attribute difference between ego and alter.

Definition at line 904 of file network.hpp.

5.3.2.2 counter_ctriads() [1/2]

Definition at line 665 of file network.hpp.

5.3.2.3 counter ctriads() [2/2]

Definition at line 610 of file network.hpp.

5.3.2.4 counter_degree()

Counts number of vertices with a given out-degree.

Definition at line 1322 of file network.hpp.

5.3.2.5 counter_density()

Definition at line 731 of file network.hpp.

5.3.2.6 counter diff()

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

Definition at line 949 of file network.hpp.

5.3 Network counters 15

5.3.2.7 counter_edges()

Number of edges.

Definition at line 152 of file network.hpp.

5.3.2.8 counter_idegree() [1/2]

Definition at line 1166 of file network.hpp.

5.3.2.9 counter_idegree() [2/2]

Counts number of vertices with a given in-degree.

Definition at line 1119 of file network.hpp.

5.3.2.10 counter_idegree15() [1/2]

Definition at line 787 of file network.hpp.

5.3.2.11 counter_idegree15() [2/2]

Definition at line 759 of file network.hpp.

5.3.2.12 counter_isolates() [1/2]

Definition at line 215 of file network.hpp.

5.3.2.13 counter_isolates() [2/2]

Number of isolated vertices.

Definition at line 175 of file network.hpp.

5.3.2.14 counter_istar2() [1/2]

Definition at line 338 of file network.hpp.

5.3.2.15 counter_istar2() [2/2]

Definition at line 312 of file network.hpp.

5.3.2.16 counter_mutual()

Number of mutual ties.

Definition at line 256 of file network.hpp.

5.3 Network counters 17

5.3.2.17 counter_nodecov()

Definition at line 1062 of file network.hpp.

5.3.2.18 counter_nodeicov()

Definition at line 1012 of file network.hpp.

5.3.2.19 counter_nodematch()

Definition at line 1087 of file network.hpp.

5.3.2.20 counter_nodeocov()

Definition at line 1037 of file network.hpp.

5.3.2.21 counter_odegree() [1/2]

Definition at line 1267 of file network.hpp.

5.3.2.22 counter_odegree() [2/2]

Counts number of vertices with a given out-degree.

Definition at line 1219 of file network.hpp.

5.3.2.23 counter_odegree15() [1/2]

Definition at line 861 of file network.hpp.

5.3.2.24 counter_odegree15() [2/2]

Definition at line 833 of file network.hpp.

5.3.2.25 counter_ostar2() [1/2]

Definition at line 404 of file network.hpp.

5.3.2.26 counter_ostar2() [2/2]

Definition at line 376 of file network.hpp.

5.4 Phylo counters 19

5.3.2.27 counter_ttriads() [1/2]

Definition at line 531 of file network.hpp.

5.3.2.28 counter_ttriads() [2/2]

Definition at line 441 of file network.hpp.

5.3.2.29 NETWORK_COUNTER()

Definition at line 993 of file network.hpp.

5.4 Phylo counters

Counters for phylogenetic modeling.

Functions

- void counter_overall_gains (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)
 Overall functional gains.
- void counter_gains (PhyloCounters *counters, std::vector< uint > nfun, unsigned int duplication=DEFAULT_DUPLICATION)

 Functional gains for a specific function (nfun).
- void counter_gains_k_offspring (PhyloCounters *counters, std::vector< uint > nfun, uint k=1u, unsigned int duplication=DEFAULT_DUPLICATION)

k genes gain function nfun

- void counter_genes_changing (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)
- void counter_preserve_pseudogene (PhyloCounters *counters, unsigned int nfunA, unsigned int nfunB, unsigned int duplication=DEFAULT DUPLICATION)

Keeps track of how many pairs of genes preserve pseudostate.

- void counter_prop_genes_changing (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)
- void counter_overall_loss (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

Overall functional loss.

• void counter_maxfuns (PhyloCounters *counters, uint lb, uint ub, unsigned int duplication=DEFAULT_DUPLICATION)

Cap the number of functions per gene.

- void counter_loss (PhyloCounters *counters, std::vector< uint > nfun, unsigned int duplication=DEFAULT_DUPLICATION)

 Total count of losses for an specific function.
- void counter_overall_changes (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

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

 Total count of Sub-functionalization events.
- void counter_cogain (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

 Co-evolution (joint gain or loss)
- void counter_longest (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Longest branch mutates (either by gain or by loss)
- void counter_neofun (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

 Total number of neofunctionalization events.
- void counter_pairwise_neofun_singlefun (PhyloCounters *counters, uint nfunA, unsigned int duplication=DEFAULT_DUPLICATI
 Total number of neofunctionalization events sum_u sum_{w < u} [x(u,a)*(1 x(w,a)) + (1 x(u,a)) * x(w,a)] change</p>
- stat: delta{x(u,a): 0->1} = 1 2 * x(w,a)
 void counter_neofun_a2b (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION
 Total number of neofunctionalization events.
- void counter_co_opt (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

 Function co-opting.
- void counter_k_genes_changing (PhyloCounters *counters, unsigned int k, unsigned int duplication=DEFAULT_DUPLICATION)
- void counter_less_than_p_prop_genes_changing (PhyloCounters *counters, double p, unsigned int duplication=DEFAULT_DUPLICATION)

Indicator function. Equals to one if k genes changed and zero otherwise.

Indicator function. Equals to one if k genes changed and zero otherwise.

- void counter_gains_from_0 (PhyloCounters *counters, std::vector < uint > nfun, unsigned int duplication=DEFAULT_DUPLICAT
 Used when all the functions are in 0 (like the root node prob.)
- void counter_overall_gains_from_0 (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Used when all the functions are in 0 (like the root node prob.)
- void counter_pairwise_overall_change (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Used when all the functions are in 0 (like the root node prob.)
- void counter_pairwise_preserving (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

Used when all the functions are in 0 (like the root node prob.)

 void counter_pairwise_first_gain (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

Used when all the functions are in 0 (like the root node prob.)

5.4.1 Detailed Description

Counters for phylogenetic modeling.

Parameters

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

5.4 Phylo counters 21

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 1393 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 888 of file phylo.hpp.

5.4.2.3 counter_gains()

Functional gains for a specific function (nfun).

Definition at line 193 of file phylo.hpp.

5.4.2.4 counter_gains_from_0()

Used when all the functions are in 0 (like the root node prob.)

Needs to specify function a.

Definition at line 1727 of file phylo.hpp.

5.4.2.5 counter_gains_k_offspring()

k genes gain function nfun

Definition at line 253 of file phylo.hpp.

5.4.2.6 counter_genes_changing()

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

Definition at line 325 of file phylo.hpp.

5.4.2.7 counter_k_genes_changing()

Indicator function. Equals to one if k genes changed and zero otherwise.

Definition at line 1491 of file phylo.hpp.

5.4 Phylo counters 23

5.4.2.8 counter_less_than_p_prop_genes_changing()

Indicator function. Equals to one if k genes changed and zero otherwise.

< How many genes diverge the parent

Definition at line 1611 of file phylo.hpp.

5.4.2.9 counter_longest()

Longest branch mutates (either by gain or by loss)

Definition at line 945 of file phylo.hpp.

5.4.2.10 counter_loss()

Total count of losses for an specific function.

Definition at line 688 of file phylo.hpp.

5.4.2.11 counter_maxfuns()

Cap the number of functions per gene.

Definition at line 626 of file phylo.hpp.

5.4.2.12 counter_neofun()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1115 of file phylo.hpp.

5.4.2.13 counter_neofun_a2b()

Total number of neofunctionalization events.

Needs to specify pairs of function.

Definition at line 1260 of file phylo.hpp.

5.4.2.14 counter_overall_changes()

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

Definition at line 740 of file phylo.hpp.

5.4.2.15 counter overall gains()

Overall functional gains.

Total number of gains (irrespective of the function).

Definition at line 155 of file phylo.hpp.

5.4 Phylo counters 25

5.4.2.16 counter_overall_gains_from_0()

Used when all the functions are in 0 (like the root node prob.)

Needs to specify function a.

Definition at line 1793 of file phylo.hpp.

5.4.2.17 counter_overall_loss()

Overall functional loss.

Definition at line 578 of file phylo.hpp.

5.4.2.18 counter pairwise first gain()

Used when all the functions are in 0 (like the root node prob.)

Needs to specify function a. sum $x(a)^3(1-x(b))^3 + x(b)^3(1-x(a))^3 + x(a)^3 + x(b)^3 + (1-x(a))^3 + (1-x($

Definition at line 2045 of file phylo.hpp.

5.4.2.19 counter pairwise neofun singlefun()

Total number of neofunctionalization events sum_u sum_ $\{w < u\} [x(u,a)*(1 - x(w,a)) + (1 - x(u,a)) * x(w,a)] change stat: delta<math>\{x(u,a): 0->1\} = 1 - 2 * x(w,a)$

Definition at line 1196 of file phylo.hpp.

5.4.2.20 counter_pairwise_overall_change()

Used when all the functions are in 0 (like the root node prob.)

Needs to specify function a.

Definition at line 1841 of file phylo.hpp.

5.4.2.21 counter pairwise preserving()

Used when all the functions are in 0 (like the root node prob.)

Needs to specify function a. sum $x(a)^3(1-x(b))^3 + x(b)^3(1-x(a))^3 + x(a)^3 + x(b)^3 + (1-x(a))^3 + (1-x(b))^3$ Definition at line 1906 of file phylo.hpp.

5.4.2.22 counter_preserve_pseudogene()

Keeps track of how many pairs of genes preserve pseudostate.

Definition at line 394 of file phylo.hpp.

5.4.2.23 counter_prop_genes_changing()

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

Definition at line 476 of file phylo.hpp.

5.5 Phylo rules 27

5.4.2.24 counter_subfun()

Total count of Sub-functionalization events.

It requires to specify data = {funA, funB}

Definition at line 799 of file phylo.hpp.

5.5 Phylo rules

Rules for phylogenetic modeling.

Classes

· class PhyloRuleDynData

Functions

void rule_dyn_limit_changes (PhyloSupport *support, uint pos, uint lb, uint ub, unsigned int duplication=DEFAULT_DUPLICATIC
 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()

```
uint lb, uint ub, unsigned int duplication = DEFAULT\_DUPLICATION ) [inline]
```

Overall functional gains.

Parameters

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

Returns

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

Definition at line 2175 of file phylo.hpp.

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 28 of file typedefs.hpp.

6.5.2.2 NONE

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

Definition at line 29 of file typedefs.hpp.

6.5.2.3 ONE

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

Definition at line 30 of file typedefs.hpp.

6.5.2.4 TWO

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

Definition at line 31 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 46 of file typedefs.hpp.

6.6.2.2 AS_ZERO

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

Definition at line 45 of file typedefs.hpp.

6.6.2.3 BOTH

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

Definition at line 39 of file typedefs.hpp.

6.6.2.4 NONE

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

Definition at line 40 of file typedefs.hpp.

6.6.2.5 ONE

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

Definition at line 41 of file typedefs.hpp.

6.6.2.6 TWO

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

Definition at line 42 of file typedefs.hpp.

6.6.2.7 UKNOWN

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

Definition at line 44 of file typedefs.hpp.

Class Documentation

7.1 BArray < Cell_Type, Data_Type > Class Template Reference

Baseline class for binary arrays.

#include <barray-bones.hpp>

Public Member Functions

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

Get the edgelist.

- void transpose ()
- void clear (bool hard=true)
- void resize (uint N_, uint M_)
- void reserve ()
- void print (const char *fmt=nullptr,...) const
- bool is_dense () const noexcept

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.

34 Class Documentation

• BArray ()

Zero-size array.

BArray (uint N_, uint M_)

Empty array.

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

Edgelist with data.

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

Edgelist with no data (simpler)

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

Move operator.

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

Set the data object.

- Data_Type * D ()
- const Data_Type * D () const
- · void flush_data ()

Queries

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

Parameters

i,j	Coordinates
check_bounds	If false avoids checking bounds.

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

Cell-wise insertion/deletion

Parameters

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

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

- BArray< Cell_Type, Data_Type > & operator-= (const std::pair< uint, uint > &coords)
- BArrayCell< Cell_Type, Data_Type > operator() (uint i, uint j, bool check_bounds=true)
- const Cell_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 63 of file barray-bones.hpp.

7.1.2.2 BArray() [2/6]

Empty array.

Definition at line 66 of file barray-bones.hpp.

7.1.2.3 BArray() [3/6]

Edgelist with data.

7.1.2.4 BArray() [4/6]

Edgelist with no data (simpler)

7.1.2.5 BArray() [5/6]

Copy constructor.

7.1.2.6 BArray() [6/6]

Move operator.

7.1.2.7 \sim BArray()

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

7.1.3 Member Function Documentation

7.1.3.1 clear()

7.1.3.2 col()

7.1.3.3 D() [1/2]

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

7.1.3.4 D() [2/2]

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

7.1.3.5 default_val()

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

7.1.3.6 flush_data()

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

7.1.3.7 get_cell()

7.1.3.8 get_col_vec() [1/2]

7.1.3.9 get_col_vec() [2/2]

7.1.3.10 get entries()

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

Get the edgelist.

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

Returns

Entries<Cell_Type>

7.1.3.11 get_row_vec() [1/2]

7.1.3.12 get_row_vec() [2/2]

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

7.1.3.13 insert_cell() [1/3]

7.1.3.14 insert_cell() [2/3]

7.1.3.15 insert_cell() [3/3]

7.1.3.16 is_dense()

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

Definition at line 232 of file barray-bones.hpp.

7.1.3.17 is_empty()

7.1.3.18 ncol()

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

7.1.3.19 nnozero()

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

7.1.3.20 nrow()

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

7.1.3.21 operator()() [1/2]

7.1.3.22 operator()() [2/2]

7.1.3.23 operator*=()

7.1.3.24 operator+=() [1/3]

7.1.3.25 operator+=() [2/3]

7.1.3.26 operator+=() [3/3]

7.1.3.27 operator-=() [1/3]

7.1.3.28 operator-=() [2/3]

7.1.3.29 operator-=() [3/3]

7.1.3.30 operator/=()

7.1.3.31 operator=() [1/2]

Move assignment.

7.1.3.32 operator=() [2/2]

Assignment constructor.

7.1.3.33 operator==()

7.1.3.34 out_of_range()

7.1.3.35 print()

7.1.3.36 reserve()

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

7.1.3.37 resize()

7.1.3.38 rm_cell()

7.1.3.39 row()

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

7.1.3.40 set_data()

Set the data object.

Parameters

data_	
delete_ <i>←</i>	
data_	

7.1.3.41 swap_cells()

7.1.3.42 swap_cols()

7.1.3.43 swap_rows()

7.1.3.44 toggle_cell()

7.1.3.45 toggle_lock()

7.1.3.46 transpose()

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

7.1.3.47 zero_col()

7.1.3.48 zero_row()

7.1.4 Friends And Related Function Documentation

${\bf 7.1.4.1 \quad BArrayCell} < {\bf Cell_Type, Data_Type} >$

```
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 48 of file barray-bones.hpp.

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

• include/barry/barray-bones.hpp

7.2 BArrayCell< Cell_Type, Data_Type > Class Template Reference

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

Public Member Functions

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

7.2.1 Detailed Description

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

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

7.2.2 Constructor & Destructor Documentation

7.2.2.1 BArrayCell()

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

7.2.2.2 ∼BArrayCell()

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

Definition at line 31 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
- include/barry/barrayrow-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 46 of file barraycell-bones.hpp.

7.3.2 Constructor & Destructor Documentation

7.3.2.1 BArrayCell const()

Definition at line 55 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 67 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
- · include/barry/barrayrow-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
- $\bullet \ \ \mathsf{void} \ \mathsf{get_row_vec} \ (\mathsf{std}:: \mathsf{vector} < \mathsf{Cell_Type} > *\mathsf{x}, \ \mathsf{uint} \ \mathsf{i}, \ \mathsf{bool} \ \mathsf{check_bounds} = \mathsf{true}) \ \mathsf{const}$
- BArrayDenseRow< Cell_Type, Data_Type > & row (uint i, bool check_bounds=true)
- const BArrayDenseRow_const< Cell_Type, Data_Type > row (uint i, bool check_bounds=true) const
- BArrayDenseCol< Cell_Type, Data_Type > & col (uint j, bool check_bounds=true)
- const BArrayDenseCol_const< Cell_Type, Data_Type > col (uint j, bool check_bounds=true) const
- Entries < Cell_Type > get_entries () const

Get the edgelist.

- void transpose ()
- void clear (bool hard=true)
- void resize (uint N , uint M)
- · void reserve ()
- void print (const char *fmt=nullptr,...) const
- · bool is dense () const noexcept
- const std::vector< Cell_Type > & get_data () const
- const Cell_Type rowsum (unsigned int i) const
- const Cell_Type colsum (unsigned int i) const

Constructors

Parameters

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

• BArrayDense ()

Zero-size array.

BArrayDense (uint N_, uint M_)

Empty array.

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

Edgelist with data.

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

Edgelist with no data (simpler)

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

Assignment constructor.

- $\bullet \ \ \mathsf{BArrayDense} < \mathsf{Cell_Type}, \ \mathsf{Data_Type} > \& \mathsf{x}) \ \mathsf{noexcept} \\$
 - Move operator.
- BArrayDense< Cell_Type, Data_Type > & operator= (BArrayDense< Cell_Type, Data_Type > &&x) noexcept

Move assignment.

void set_data (Data_Type *data_, bool delete_data_=false)

Set the data object.

- Data_Type * D ()
- const Data_Type * D () const

Queries

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

Parameters

i,j	Coordinates
check_bounds	If false avoids checking bounds.

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

Cell-wise insertion/deletion

Parameters

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

- BArrayDense< Cell_Type, Data_Type > & operator+= (const std::pair< uint, uint > &coords)
- BArrayDense < Cell_Type, Data_Type > & operator = (const std::pair < uint, uint > &coords)
- BArrayDenseCell< Cell_Type, Data_Type > operator() (uint i, uint j, bool check_bounds=true)
- const Cell_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 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 BArrayDenseCell
 Cell_Type, Data_Type
- class BArrayDenseCol< Cell_Type, Data_Type >
- class BArrayDenseCol_const< Cell_Type, Data_Type >
- class BArrayDenseRow< Cell Type, Data Type >
- class BArrayDenseRow_const< Cell_Type, Data_Type >

7.4.1 Detailed Description

 $\label{template} $$ \ensuremath{\sf template}$$ $$ \ensuremath{\sf cell_Type}$ = bool, typename Data_Type = bool> $$ \ensuremath{\sf 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 34 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 79 of file barraydense-bones.hpp.

7.4.2.2 BArrayDense() [2/6]

Empty array.

Definition at line 82 of file barraydense-bones.hpp.

7.4.2.3 BArrayDense() [3/6]

Edgelist with data.

7.4.2.4 BArrayDense() [4/6]

Edgelist with no data (simpler)

7.4.2.5 BArrayDense() [5/6]

Copy constructor.

7.4.2.6 BArrayDense() [6/6]

Move operator.

7.4.2.7 ∼BArrayDense()

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

7.4.3 Member Function Documentation

7.4.3.1 clear()

7.4.3.2 col() [1/2]

Definition at line 482 of file barraydense-meat.hpp.

7.4.3.3 col() [2/2]

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

7.4.3.4 colsum()

7.4.3.5 D() [1/2]

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

7.4.3.6 D() [2/2]

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

7.4.3.7 default_val()

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

7.4.3.8 get_cell()

7.4.3.9 get_col_vec() [1/2]

7.4.3.10 get_col_vec() [2/2]

7.4.3.11 get_data()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
const std::vector< Cell_Type >& BArrayDense< Cell_Type, Data_Type >::get_data ( ) const
```

7.4.3.12 get_entries()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Entries<Cell_Type> BArrayDense< 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>
```

7.4.3.13 get_row_vec() [1/2]

7.4.3.14 get_row_vec() [2/2]

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

7.4.3.15 insert_cell() [1/2]

7.4.3.16 insert cell() [2/2]

7.4.3.17 is_dense()

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

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

7.4.3.18 is_empty()

7.4.3.19 ncol()

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

7.4.3.20 nnozero()

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

7.4.3.21 nrow()

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

7.4.3.22 operator()() [1/2]

7.4.3.23 operator()() [2/2]

7.4.3.24 operator*=()

7.4.3.25 operator+=() [1/3]

7.4.3.26 operator+=() [2/3]

7.4.3.27 operator+=() [3/3]

7.4.3.28 operator-=() [1/3]

7.4.3.29 operator-=() [2/3]

7.4.3.30 operator-=() [3/3]

7.4.3.31 operator/=()

7.4.3.32 operator=() [1/2]

Move assignment.

7.4.3.33 operator=() [2/2]

Assignment constructor.

7.4.3.34 operator==()

7.4.3.35 out_of_range()

7.4.3.36 print()

7.4.3.37 reserve()

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

7.4.3.38 resize()

7.4.3.39 rm_cell()

7.4.3.40 row() [1/2]

7.4.3.41 row() [2/2]

7.4.3.42 rowsum()

7.4.3.43 set_data()

Set the data object.

Parameters

data_	
delete_ <i>←</i>	
data_	

7.4.3.44 swap_cells()

7.4.3.45 swap_cols()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
void BArrayDense< Cell_Type, Data_Type >::swap_cols (
         uint j0,
         uint j1,
         bool check_bounds = true )
```

7.4.3.46 swap_rows()

7.4.3.47 toggle_cell()

7.4.3.48 toggle_lock()

7.4.3.49 transpose()

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

7.4.3.50 zero_col()

7.4.3.51 zero_row()

7.4.4 Friends And Related Function Documentation

7.4.4.1 BArrayDenseCell< Cell_Type, Data_Type >

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

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

7.4.4.2 BArrayDenseCol < Cell_Type, Data_Type >

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

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

7.4.4.3 BArrayDenseCol_const< Cell_Type, Data_Type >

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

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

7.4.4.4 BArrayDenseRow < Cell_Type, Data_Type >

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

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

7.4.4.5 BArrayDenseRow_const< Cell_Type, Data_Type >

```
template<typename Cell_Type = bool, typename Data_Type = bool>
friend class BArrayDenseRow_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 64 of file barraydense-bones.hpp.

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

- · include/barry/barraydense-bones.hpp
- include/barry/barraydense-meat.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 i_, 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

Friends

- class BArrayDense< Cell_Type, Data_Type >
- class BArrayDenseCol< Cell Type, Data Type >
- class BArrayDenseCol_const< Cell_Type, Data_Type >

7.5.1 Detailed Description

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

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

7.5.2 Constructor & Destructor Documentation

7.5.2.1 BArrayDenseCell()

Definition at line 27 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 49 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 57 of file barraydensecell-meat.hpp.

7.5.3.2 operator*=()

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

7.5.3.3 operator+=()

Definition at line 19 of file barraydensecell-meat.hpp.

7.5.3.4 operator-=()

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

7.5.3.5 operator/=()

Definition at line 47 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 62 of file barraydensecell-meat.hpp.

7.5.4 Friends And Related Function Documentation

7.5.4.1 BArrayDense < Cell_Type, Data_Type >

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

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

7.5.4.2 BArrayDenseCol < Cell_Type, Data_Type >

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

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

7.5.4.3 BArrayDenseCol_const< Cell_Type, Data_Type >

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

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

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

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

7.6 BArrayDenseCell_const< Cell_Type, Data_Type > Class Template Reference

7.6.1 Detailed Description

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

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

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

• include/barry/barraydense-bones.hpp

7.7 BArrayDenseCol< Cell_Type, Data_Type > Class Template Reference

#include <barraydensecol-bones.hpp>

Public Member Functions

- BArrayDenseCol (BArrayDense < Cell_Type, Data_Type > & array_, unsigned int j)
- Col_type< Cell_Type >::iterator & begin ()
- Col_type< Cell_Type >::iterator & end ()
- size_t size () const noexcept
- std::pair< unsigned int, Cell_Type * > & operator() (unsigned int i)

Friends

- class BArrayDense< Cell_Type, Data_Type >
- class BArrayDenseCell
 Cell Type, Data Type
- class BArrayDenseCell_const< Cell_Type, Data_Type >

7.7.1 Detailed Description

```
\label{template} \mbox{typename Cell_Type = bool, typename Data_Type = bool} > \mbox{class BArrayDenseCol} < \mbox{Cell_Type, Data_Type} > \mbox{}
```

Definition at line 9 of file barraydensecol-bones.hpp.

7.7.2 Constructor & Destructor Documentation

7.7.2.1 BArrayDenseCol()

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

7.7.3 Member Function Documentation

7.7.3.1 begin()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Col_type<Cell_Type>::iterator& BArrayDenseCol< Cell_Type, Data_Type >::begin ( ) [inline]
```

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

7.7.3.2 end()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Col_type<Cell_Type>::iterator& BArrayDenseCol< Cell_Type, Data_Type >::end ( ) [inline]
```

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

7.7.3.3 operator()()

```
template<typename Cell_Type = bool, typename Data_Type = bool> std::pair<unsigned int,Cell_Type*>& BArrayDenseCol< Cell_Type, Data_Type >::operator() ( unsigned int i ) [inline]
```

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

7.7.3.4 size()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
size_t BArrayDenseCol< Cell_Type, Data_Type >::size ( ) const [inline], [noexcept]
```

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

7.7.4 Friends And Related Function Documentation

7.7.4.1 BArrayDense < Cell_Type, Data_Type >

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

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

7.7.4.2 BArrayDenseCell< Cell_Type, Data_Type >

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

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

7.7.4.3 BArrayDenseCell_const< Cell_Type, Data_Type >

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

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

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

- include/barry/barraydense-meat.hpp
- include/barry/barraydensecol-bones.hpp

7.8 BArrayDenseCol_const< Cell_Type, Data_Type > Class Template Reference

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

Public Member Functions

- BArrayDenseCol_const (const BArrayDense< Cell_Type, Data_Type > &array_, unsigned int j)
- Col_type< Cell_Type >::iterator begin ()
- Col_type< Cell_Type >::iterator end ()
- size t size () const noexcept
- const std::pair< unsigned int, Cell_Type * > operator() (unsigned int i) const

Friends

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

7.8.1 Detailed Description

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

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

7.8.2 Constructor & Destructor Documentation

7.8.2.1 BArrayDenseCol_const()

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

7.8.3 Member Function Documentation

7.8.3.1 begin()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Col_type<Cell_Type>::iterator BArrayDenseCol_const< Cell_Type, Data_Type >::begin ( ) [inline]
```

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

7.8.3.2 end()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Col_type<Cell_Type>::iterator BArrayDenseCol_const< Cell_Type, Data_Type >::end ( ) [inline]
```

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

7.8.3.3 operator()()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
const std::pair<unsigned int,Cell_Type*> BArrayDenseCol_const< Cell_Type, Data_Type >::operator()
(
    unsigned int i ) const [inline]
```

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

7.8.3.4 size()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
size_t BArrayDenseCol_const< Cell_Type, Data_Type >::size ( ) const [inline], [noexcept]
```

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

7.8.4 Friends And Related Function Documentation

7.8.4.1 BArrayDenseCell< Cell_Type, Data_Type >

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

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

7.8.4.2 BArrayDenseCell_const< Cell_Type, Data_Type >

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

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

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

- include/barry/barraydense-meat.hpp
- include/barry/barraydensecol-bones.hpp

7.9 BArrayDenseRow< Cell_Type, Data_Type > Class Template Reference

#include <barraydenserow-bones.hpp>

Public Member Functions

- BArrayDenseRow (BArrayDense < Cell_Type, Data_Type > & array_, unsigned int i)
- Row_type< Cell_Type >::iterator & begin ()
- Row_type< Cell_Type >::iterator & end ()
- size_t size () const noexcept
- std::pair< unsigned int, Cell< Cell_Type > > & operator() (unsigned int i)

Friends

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

7.9.1 Detailed Description

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

Definition at line 9 of file barraydenserow-bones.hpp.

7.9.2 Constructor & Destructor Documentation

7.9.2.1 BArrayDenseRow()

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

7.9.3 Member Function Documentation

7.9.3.1 begin()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Row_type<Cell_Type>::iterator& BArrayDenseRow< Cell_Type, Data_Type >::begin ( ) [inline]
```

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

7.9.3.2 end()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Row_type<Cell_Type>::iterator& BArrayDenseRow< Cell_Type, Data_Type >::end ( ) [inline]
```

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

7.9.3.3 operator()()

```
\label{template} $$ \text{template}$ $$ \text{typename Cell_Type = bool, typename Data_Type = bool} $$ \text{std::pair}$ \text{cunsigned int,Cell}$ $$ \text{Cell_Type}$ > & $$ BArrayDenseRow$ Cell_Type, Data_Type >::operator() $$ ($$ unsigned int $i$ ) [inline] $$
```

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

7.9.3.4 size()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
size_t BArrayDenseRow< Cell_Type, Data_Type >::size ( ) const [inline], [noexcept]
```

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

7.9.4 Friends And Related Function Documentation

7.9.4.1 BArrayDense< Cell_Type, Data_Type>

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

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

7.9.4.2 BArrayDenseCell< Cell_Type, Data_Type >

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

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

7.9.4.3 BArrayDenseCell_const< Cell_Type, Data_Type >

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

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

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

- include/barry/barraydense-bones.hpp
- include/barry/barraydenserow-bones.hpp

7.10 BArrayDenseRow_const< Cell_Type, Data_Type > Class Template Reference

#include <barraydenserow-bones.hpp>

Public Member Functions

- BArrayDenseRow_const (const BArrayDense< Cell_Type, Data_Type > &array_, unsigned int i)
- Row_type< Cell_Type >::const_iterator begin () const
- Row_type< Cell_Type >::const_iterator end () const
- size_t size () const noexcept
- const std::pair< unsigned int, Cell< Cell_Type >> operator() (unsigned int i) const

Friends

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

7.10.1 Detailed Description

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

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

7.10.2 Constructor & Destructor Documentation

7.10.2.1 BArrayDenseRow_const()

Definition at line 89 of file barraydenserow-bones.hpp.

7.10.3 Member Function Documentation

7.10.3.1 begin()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Row_type< Cell_Type >::const_iterator BArrayDenseRow_const< Cell_Type, Data_Type >::begin ( )
const [inline]
```

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

7.10.3.2 end()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
Row_type< Cell_Type >::const_iterator BArrayDenseRow_const< Cell_Type, Data_Type >::end ( )
const [inline]
```

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

7.10.3.3 operator()()

Definition at line 123 of file barraydenserow-bones.hpp.

7.10.3.4 size()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
size_t BArrayDenseRow_const< Cell_Type, Data_Type >::size ( ) const [inline], [noexcept]
```

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

7.10.4 Friends And Related Function Documentation

7.10.4.1 BArrayDenseCell< Cell $_$ Type, Data $_$ Type>

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

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

7.10.4.2 BArrayDenseCell_const < Cell_Type, Data_Type >

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

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

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

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

7.11 BArrayRow < Cell_Type, Data_Type > Class Template Reference

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

Public Member Functions

- BArrayRow (BArray< Cell_Type, Data_Type > *Array_, uint i_, bool check_bounds=true)
- ∼BArrayRow ()
- void operator= (const BArrayRow< Cell Type, Data Type > &val)
- void operator+= (const BArrayRow< Cell_Type, Data_Type > &val)
- void operator-= (const BArrayRow< Cell_Type, Data_Type > &val)
- $\bullet \ \ \mathsf{void} \ \mathsf{operator} *= (\mathsf{const} \ \mathsf{BArrayRow} < \mathsf{Cell_Type}, \ \mathsf{Data_Type} > \mathsf{\&val})$
- void operator/= (const BArrayRow< Cell_Type, Data_Type > &val)
- operator BArrayRow
 Cell_Type, Data_Type > () const
- bool operator== (const BArrayRow< Cell_Type, Data_Type > &val) const

7.11.1 Detailed Description

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

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

7.11.2 Constructor & Destructor Documentation

7.11.2.1 BArrayRow()

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

7.11.2.2 ∼BArrayRow()

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

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

7.11.3 Member Function Documentation

7.11.3.1 operator BArrayRow< Cell_Type, Data_Type >()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
BArrayRow< Cell_Type, Data_Type >::operator BArrayRow< Cell_Type, Data_Type > ( ) const
```

7.11.3.2 operator*=()

7.11.3.3 operator+=()

7.11.3.4 operator-=()

7.11.3.5 operator/=()

7.11.3.6 operator=()

7.11.3.7 operator==()

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

• include/barry/barrayrow-bones.hpp

7.12 BArrayRow_const< Cell_Type, Data_Type > Class Template Reference

#include <barrayrow-bones.hpp>

Public Member Functions

- BArrayRow_const (const BArray < Cell_Type, Data_Type > *Array_, uint i_, bool check_bounds=true)
- ∼BArrayRow_const ()
- operator BArrayRow_const< Cell_Type, Data_Type > () const
- bool operator== (const BArrayRow_const< Cell_Type, Data_Type > &val) const
- bool operator!= (const BArrayRow const< Cell Type, Data Type > &val) const
- bool operator< (const BArrayRow_const< Cell_Type, Data_Type > &val) const
- bool operator> (const BArrayRow_const< Cell_Type, Data_Type > &val) const
- bool operator<= (const BArrayRow_const< Cell_Type, Data_Type > &val) const
- bool operator>= (const BArrayRow_const< Cell_Type, Data_Type > &val) const

7.12.1 Detailed Description

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

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

7.12.2 Constructor & Destructor Documentation

7.12.2.1 BArrayRow_const()

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

7.12.2.2 ~BArrayRow_const()

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

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

7.12.3 Member Function Documentation

7.12.3.1 operator BArrayRow_const< Cell_Type, Data_Type >()

```
template<typename Cell_Type = bool, typename Data_Type = bool>
BArrayRow_const< Cell_Type, Data_Type >::operator BArrayRow_const< Cell_Type, Data_Type > ( )
const
```

7.12.3.2 operator"!=()

7.12.3.3 operator<()

7.12.3.4 operator<=()

7.12.3.5 operator==()

7.12.3.6 operator>()

7.12.3.7 operator>=()

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

• include/barry/barrayrow-bones.hpp

7.13 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

7.13.1 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.

7.13.2 Constructor & Destructor Documentation

7.13.2.1 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.

7.13.2.2 ~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.

7.13.3 Member Function Documentation

7.13.3.1 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.

7.13.3.2 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.

7.13.3.3 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.

7.13.3.4 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.

7.13.3.5 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.

7.13.3.6 operator*=()

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

7.13.3.7 operator+=()

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

7.13.3.8 operator-=()

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

7.13.3.9 operator/=()

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

7.13.3.10 operator=()

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

7.13.3.11 operator==()

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

7.13.3.12 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

7.14 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

7.14.1 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.

7.14.2 Constructor & Destructor Documentation

7.14.2.1 BArrayVector_const()

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

7.14.2.2 ~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.

7.14.3 Member Function Documentation

7.14.3.1 begin()

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

7.14.3.2 end()

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

7.14.3.3 is_col()

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

7.14.3.4 is_row()

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

7.14.3.5 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.

7.14.3.6 operator"!=()

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

7.14.3.7 operator<()

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

7.14.3.8 operator<=()

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

7.14.3.9 operator==()

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

7.14.3.10 operator>()

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

7.14.3.11 operator>=()

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

7.14.3.12 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

7.15 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, bool active_=true)
- ∼Cell ()
- Cell (const Cell
 Cell_Type > &arg)
- Cell< Cell_Type > & operator= (const 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
- bool active

7.15.1 Detailed Description

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

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

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

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

7.15.2 Constructor & Destructor Documentation

7.15.2.1 Cell() [1/7]

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

7.15.2.2 Cell() [2/7]

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

7.15.2.3 ∼Cell()

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

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

7.15.2.4 Cell() [3/7]

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

7.15.2.5 Cell() [4/7]

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

7.15.2.6 Cell() [5/7]

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

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

7.15.2.7 Cell() [6/7]

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

Definition at line 65 of file cell-meat.hpp.

7.15.2.8 Cell() [7/7]

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

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

7.15.3 Member Function Documentation

7.15.3.1 add() [1/4]

7.15.3.2 add() [2/4]

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

7.15.3.3 add() [3/4]

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

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

7.15.3.4 add() [4/4]

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

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

7.15.3.5 operator Cell_Type()

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

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

7.15.3.6 operator"!=()

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

7.15.3.7 operator=() [1/2]

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

7.15.3.8 operator=() [2/2]

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

7.15.3.9 operator==()

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

7.15.4 Member Data Documentation

7.15.4.1 active

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

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

7.15.4.2 value

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

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

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

7.16 Cell_const < Cell_Type > Class Template Reference

7.16.1 Detailed Description

```
template<typename Cell_Type> class Cell_const< Cell_Type>
```

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

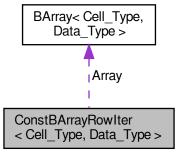
The documentation for this class was generated from the following file:

• include/barry/barray-meat.hpp

7.17 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

7.17.1 Detailed Description

```
\label{template} \begin{tabular}{ll} template < typename Cell_Type, typename Data_Type > \\ class ConstBArrayRowlter < Cell_Type, Data_Type > \\ \end{tabular}
```

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

7.17.2 Constructor & Destructor Documentation

7.17.2.1 ConstBArrayRowIter()

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

7.17.2.2 ~ConstBArrayRowlter()

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

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

7.17.3 Member Data Documentation

7.17.3.1 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.

7.17.3.2 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.

7.17.3.3 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.

7.17.3.4 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

7.18 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)
- std::string get_name () const
- std::string get_description () const

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_, Data_Type data_, 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
- std::string name = ""
- std::string desc = ""

7.18.1 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.

7.18.2 Constructor & Destructor Documentation

7.18.2.1 Counter() [1/4]

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

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

7.18.2.2 Counter() [2/4]

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

7.18.2.3 Counter() [3/4]

Copy constructor.

7.18.2.4 Counter() [4/4]

Move constructor.

7.18.2.5 ∼Counter()

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

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

7.18.3 Member Function Documentation

7.18.3.1 count()

7.18.3.2 get_description()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
std::string Counter< Array_Type, Data_Type >::get_description ( ) const
```

7.18.3.3 get_name()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
std::string Counter< Array_Type, Data_Type >::get_name ( ) const
```

7.18.3.4 init()

7.18.3.5 operator=() [1/2]

Copy assignment.

7.18.3.6 operator=() [2/2]

Move assignment.

7.18.4 Member Data Documentation

7.18.4.1 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.

7.18.4.2 data

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

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

7.18.4.3 desc

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

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

7.18.4.4 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.

7.18.4.5 name

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

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

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

• include/barry/counters-bones.hpp

7.19 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_fun_type< Array_Type, Data_Type > count_fun_, Counter_fun_type< Array_

 Type, Data_Type > init_fun_, Data_Type data_, std::string name_="", std::string desc_="")
- std::vector< std::string > get_names () const
- std::vector< std::string > get_descriptions () const

7.19.1 Detailed Description

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

Vector of counters.

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

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

7.19.2 Constructor & Destructor Documentation

7.19.2.1 Counters() [1/3]

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

7.19.2.2 ∼Counters()

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

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

7.19.2.3 Counters() [2/3]

Copy constructor.

Parameters



7.19.2.4 Counters() [3/3]

Move constructor.

Parameters

counters⊷

7.19.3 Member Function Documentation

7.19.3.1 add_counter() [1/2]

7.19.3.2 add_counter() [2/2]

7.19.3.3 get_descriptions()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
std::vector< std::string > Counters< Array_Type, Data_Type >::get_descriptions ( ) const
```

7.19.3.4 get_names()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
std::vector< std::string > Counters< Array_Type, Data_Type >::get_names ( ) const
```

7.19.3.5 operator=() [1/2]

Copy assignment constructor.

Parameters

counter⊷	

Returns

Counters<Array_Type,Data_Type>

7.19.3.6 operator=() [2/2]

Move assignment constructor.

Parameters



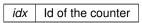
Returns

Counters<Array_Type,Data_Type>&

7.19.3.7 operator[]()

Returns a pointer to a particular counter.

Parameters



Returns

Counter<Array_Type,Data_Type>*

7.19.3.8 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 150 of file counters-bones.hpp.

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

• include/barry/counters-bones.hpp

7.20 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
- $std::vector < Cell_Type > val$

7.20.1 Detailed Description

```
\label{template} \mbox{template}{<}\mbox{typename Cell\_Type}{>} \\ \mbox{class Entries}{<}\mbox{Cell\_Type}{>} \\
```

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

Template Parameters

Cell_Type Any type

Definition at line 79 of file typedefs.hpp.

7.20.2 Constructor & Destructor Documentation

7.20.2.1 Entries() [1/2]

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

Definition at line 85 of file typedefs.hpp.

7.20.2.2 Entries() [2/2]

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

Definition at line 86 of file typedefs.hpp.

7.20.2.3 ∼Entries()

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

Definition at line 93 of file typedefs.hpp.

7.20.3 Member Function Documentation

7.20.3.1 resize()

Definition at line 95 of file typedefs.hpp.

7.20.4 Member Data Documentation

7.20.4.1 source

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

Definition at line 81 of file typedefs.hpp.

7.20.4.2 target

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

Definition at line 82 of file typedefs.hpp.

7.20.4.3 val

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

Definition at line 83 of file typedefs.hpp.

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

• include/barry/typedefs.hpp

7.21 Flock Class Reference

A Flock is a group of Geese.

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

7.21 Flock Class Reference 111

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 (unsigned int bar width=BARRY PROGRESS BAR WIDTH)
- phylocounters::PhyloCounters * get_counters ()
- phylocounters::PhyloSupport * get support fun ()
- std::vector< std::vector< double > > * get_stats_support ()
- std::vector< std::vector< double >> * get_stats_target ()
- phylocounters::PhyloModel * get model ()

Returns the joint likelihood of the model.

• Geese * operator() (unsigned int i, bool check bounds=true)

Access the i-th geese element.

Information about the model

- · unsigned int nfuns () const noexcept
- unsigned int ntrees () const noexcept
- std::vector< unsigned int > nnodes () const noexcept
- std::vector< unsigned int > nleafs () const noexcept
- unsigned int nterms () const
- unsigned int support_size () const noexcept
- std::vector< std::string > colnames () const
- unsigned int parse_polytomies (bool verb=true, std::vector < size_t > *dist=nullptr) const noexcept
 Check polytomies and return the largest.
- · void print () const

Public Attributes

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

7.21.1 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.

7.21.2 Constructor & Destructor Documentation

7.21.2.1 Flock()

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

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

7.21.2.2 ∼Flock()

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

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

7.21.3 Member Function Documentation

7.21.3.1 add_data()

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

Add a tree to the flock.

Parameters

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

Returns

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

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

7.21 Flock Class Reference 113

7.21.3.2 colnames()

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

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

7.21.3.3 get_counters()

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

Definition at line 100 of file flock-meat.hpp.

7.21.3.4 get_model()

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

Definition at line 131 of file flock-meat.hpp.

7.21.3.5 get_stats_support()

```
std::vector< std::vector< double > > * Flock::get_stats_support ( ) [inline]
```

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

7.21.3.6 get_stats_target()

```
std::vector< std::vector< double > > * Flock::get_stats_target ( ) [inline]
```

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

7.21.3.7 get_support_fun()

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

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

7.21.3.8 init()

```
void Flock::init (
     unsigned int bar_width = BARRY_PROGRESS_BAR_WIDTH ) [inline]
```

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

7.21.3.9 likelihood_joint()

Returns the joint likelihood of the model.

Parameters

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

Returns

double

Definition at line 138 of file flock-meat.hpp.

7.21.3.10 nfuns()

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

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

7.21.3.11 nleafs()

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

Definition at line 195 of file flock-meat.hpp.

7.21 Flock Class Reference 115

7.21.3.12 nnodes()

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

Definition at line 181 of file flock-meat.hpp.

7.21.3.13 nterms()

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

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

7.21.3.14 ntrees()

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

Definition at line 174 of file flock-meat.hpp.

7.21.3.15 operator()()

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

Access the i-th geese element.

Parameters

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

Returns

Geese*

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

7.21.3.16 parse_polytomies()

```
unsigned int Flock::parse_polytomies (
          bool verb = true,
          std::vector< size_t > * dist = nullptr ) const [inline], [noexcept]
```

Check polytomies and return the largest.

Definition at line 231 of file flock-meat.hpp.

7.21.3.17 print()

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

Definition at line 258 of file flock-meat.hpp.

7.21.3.18 set_seed()

Set the seed of the model.

Parameters

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

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

7.21.3.19 support_size()

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

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

7.21.4 Member Data Documentation

7.21.4.1 dat

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

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

7.21.4.2 initialized

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

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

7.21.4.3 model

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

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

7.21.4.4 nfunctions

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

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

7.21.4.5 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-meat.hpp

7.22 FreqTable < T > Class Template Reference

Frequency table of vectors.

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

Public Member Functions

```
FreqTable ()
~FreqTable ()
size_t add (const std::vector< T > &x, size_t *h_precomp)
Counts_type as_vector () const
const std::vector< double > & get_data () const
const std::unordered_map< size_t, size_t > & get_index () const
void clear ()
void reserve (size_t n, size_t k)
void print () const
size_t size () const noexcept
Number of unique elements in the table. (.
size_t make_hash (const std::vector< double > &x) const
```

7.22.1 Detailed Description

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

Frequency table of vectors.

This is mostly used in Support. The main data is contained in the data double vector. The matrix is stored in a row-wise fashion, where the first element is the frequency with which the vector is observed.

For example, in a model with k terms the first k+1 elements of data would be:

- · weights
- term 1
- term 2
- ..
- · term k

Definition at line 27 of file statsdb.hpp.

7.22.2 Constructor & Destructor Documentation

7.22.2.1 FreqTable()

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

Definition at line 39 of file statsdb.hpp.

7.22.2.2 \sim FreqTable()

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

Definition at line 40 of file statsdb.hpp.

7.22.3 Member Function Documentation

7.22.3.1 add()

Definition at line 64 of file statsdb.hpp.

7.22.3.2 as_vector()

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

Definition at line 144 of file statsdb.hpp.

7.22.3.3 clear()

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

Definition at line 173 of file statsdb.hpp.

7.22.3.4 get_data()

```
template<typename T = double>
const std::vector< double >& FreqTable< T >::get_data ( ) const [inline]
```

Definition at line 45 of file statsdb.hpp.

7.22.3.5 get_index()

```
template<typename T = double>
const std::unordered_map<size_t,size_t>& FreqTable< T >::get_index ( ) const [inline]
```

Definition at line 46 of file statsdb.hpp.

7.22.3.6 make_hash()

Definition at line 244 of file statsdb.hpp.

7.22.3.7 print()

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

Definition at line 209 of file statsdb.hpp.

7.22.3.8 reserve()

Definition at line 187 of file statsdb.hpp.

7.22.3.9 size()

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

Number of unique elements in the table. (.

Returns

size_t

Definition at line 236 of file statsdb.hpp.

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

• include/barry/statsdb.hpp

7.23 Geese Class Reference 121

7.23 Geese Class Reference

Annotated Phylo Model.

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

Public Member Functions

- ∼Geese ()
- void init (unsigned int bar width=BARRY PROGRESS BAR WIDTH)
- void inherit_support (const Geese &model_, bool delete_support_=false)
- void calc_sequence (Node *n=nullptr)
- void calc reduced sequence ()
- double likelihood (const std::vector< double > &par, bool as_log=false, bool use_reduced_sequence=true)
- double likelihood_exhaust (const std::vector< double > &par)
- std::vector< double > get_probabilities () const
- void set_seed (const unsigned int &s)
- std::vector< std::vector< unsigned int > > simulate (const std::vector< double > &par)
- std::vector< std::vector< double > > observed counts ()
- void print_observed_counts ()
- void print () const

Prints information about the DEFM.

- void init node (Node &n)
- void update_annotations (unsigned int nodeid, std::vector< unsigned int > newann)
- $std::vector < std::vector < bool > > get_states$ () const

Powerset of a gene's possible states.

std::vector< unsigned int > get_annotated_nodes () const

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

- ∼Geese ()
- void init (unsigned int bar width=BARRY PROGRESS BAR WIDTH)
- void inherit_support (const Geese &model_, bool delete_support_=false)
- void calc_sequence (Node *n=nullptr)
- void calc reduced sequence ()
- double likelihood (const std::vector< double > &par, bool as_log=false, bool use_reduced_sequence=true)
- double likelihood_exhaust (const std::vector< double > &par)
- std::vector< double > get_probabilities () const
- void set_seed (const unsigned int &s)
- std::vector< std::vector< unsigned int > > simulate (const std::vector< double > &par)
- std::vector< std::vector< double >> observed counts ()
- void print_observed_counts ()
- · void print () const

Prints information about the GEESE.

- void init_node (Node &n)
- void update_annotations (unsigned int nodeid, std::vector< unsigned int > newann)
- std::vector< std::vector< bool > > get states () const

Powerset of a gene's possible states.

• std::vector< unsigned int > get_annotated_nodes () const

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

Construct a new Geese object

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

Parameters

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

The ordering of the entries does not matter. Passing the nodes in post order or not makes no difference to the constructor.

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

Information about the model

Parameters

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

· unsigned int nfuns () const noexcept

Number of functions analyzed.

· unsigned int nnodes () const noexcept

Number of nodes (interior + leaf)

• unsigned int nleafs () const noexcept

Number of leaf.

• unsigned int nterms () const

Number of terms included.

• unsigned int support_size () const noexcept

Number of unique sets of sufficient stats.

std::vector< unsigned int > nannotations () const noexcept

Number of annotations.

• std::vector< std::string > colnames () const

Names of the terms in the model.

unsigned int parse_polytomies (bool verb=true, std::vector< size_t > *dist=nullptr) const noexcept
 Check polytomies and return the largest.

· unsigned int nfuns () const noexcept

Number of functions analyzed.

• unsigned int nnodes () const noexcept

Number of nodes (interior + leaf)

· unsigned int nleafs () const noexcept

Number of leaf.

· unsigned int nterms () const

Number of terms included.

- unsigned int support_size () const noexcept
 - Number of unique sets of sufficient stats.
- std::vector< unsigned int > nannotations () const noexcept

Number of annotations.

- std::vector< std::string > colnames () const
 - Names of the terms in the model.
- unsigned int parse_polytomies (bool verb=true, std::vector< size_t > *dist=nullptr) 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_backend (const std::vector < double > &par, bool use_← reduced_sequence, const std::vector < uint > &preorder)
- std::vector< std::vector< double >> predict_exhaust_backend (const std::vector< double > &par, const std::vector< uint > &preorder)
- std::vector< std::vector< double > > predict_exhaust (const std::vector< double > &par)
- std::vector< std::vector< double > > predict_sim (const std::vector< double > &par, bool only_← annotated=false, unsigned int nsims=10000u)
- std::vector< std::vector< double >> predict (const std::vector< double > &par, std::vector< std::vector< double >> *res_prob=nullptr, bool leave_one_out=false, bool only_annotated=false, bool use_reduced
 _sequence=true)
- std::vector< std::vector< double > > predict_backend (const std::vector< double > &par, bool use_←
 reduced_sequence, const std::vector< uint > &preorder)
- std::vector< std::vector< double >> predict_exhaust_backend (const std::vector< double > &par, const std::vector< uint > &preorder)
- std::vector< std::vector< double > > predict_exhaust (const std::vector< double > &par)
- std::vector < std::vector < double > > predict_sim (const std::vector < double > &par, bool only_
 —
 annotated=false, unsigned int nsims=10000u)

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

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

Returns

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

std::mt19937 * get_rengine()

phylocounters::PhyloCounters * get_counters()

phylocounters::PhyloSupport * get_support_fun()

std::mt19937 * get_rengine()

phylocounters::PhyloCounters * get_counters()

phylocounters::PhyloModel * get_model()

phylocounters::PhyloModel * get_model()

phylocounters::PhyloSupport * get_support_fun()
```

Public Attributes

- · unsigned int nfunctions
- std::map< unsigned int, Node > nodes
- barry::MapVec type< unsigned int > map to nodes
- std::vector< std::vector< size t >> pset loc

Locations of columns.

- std::vector< unsigned int > sequence
- std::vector< unsigned int > reduced_sequence
- bool initialized = false
- bool delete_rengine = false
- bool delete_support = false

7.23.1 Detailed Description

Annotated Phylo Model.

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

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

7.23.2 Constructor & Destructor Documentation

7.23.2.1 Geese() [1/8]

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

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

7.23.2.2 Geese() [2/8]

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

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

7.23.2.3 Geese() [3/8]

Definition at line 214 of file geese-meat-constructors.hpp.

7.23.2.4 Geese() [4/8]

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

7.23.2.5 ~Geese() [1/2]

```
Geese::∼Geese ( ) [inline]
```

Definition at line 91 of file defm-meat.hpp.

7.23.2.6 Geese() [5/8]

```
Geese::Geese ( )
```

7.23.2.7 Geese() [6/8]

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

7.23.2.8 Geese() [7/8]

7.23.2.9 Geese() [8/8]

7.23.2.10 ~Geese() [2/2]

```
Geese::∼Geese ()
```

7.23.3 Member Function Documentation

7.23.3.1 calc_reduced_sequence() [1/2]

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

Definition at line 331 of file defm-meat.hpp.

7.23.3.2 calc_reduced_sequence() [2/2]

```
void Geese::calc_reduced_sequence ( )
```

7.23 Geese Class Reference 127

7.23.3.3 calc_sequence() [1/2]

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

Definition at line 287 of file defm-meat.hpp.

7.23.3.4 calc_sequence() [2/2]

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

7.23.3.5 colnames() [1/2]

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

Names of the terms in the model.

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

7.23.3.6 colnames() [2/2]

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

Names of the terms in the model.

7.23.3.7 get annotated nodes() [1/2]

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

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

Definition at line 669 of file defm-meat.hpp.

7.23.3.8 get_annotated_nodes() [2/2]

```
{\tt std::vector} < {\tt unsigned int} > {\tt Geese::get\_annotated\_nodes} ( ) const
```

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

7.23.3.9 get_counters() [1/2]

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

Definition at line 652 of file defm-meat.hpp.

7.23.3.10 get_counters() [2/2]

```
phylocounters::PhyloCounters* Geese::get_counters ( )
```

7.23.3.11 get_model() [1/2]

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

Definition at line 657 of file defm-meat.hpp.

7.23.3.12 get_model() [2/2]

```
phylocounters::PhyloModel* Geese::get_model ( )
```

7.23.3.13 get_probabilities() [1/2]

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

Definition at line 379 of file defm-meat.hpp.

7.23.3.14 get_probabilities() [2/2]

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

7.23.3.15 get_rengine() [1/2]

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

Definition at line 647 of file defm-meat.hpp.

7.23 Geese Class Reference 129

7.23.3.16 get_rengine() [2/2]

```
std::mt19937* Geese::get_rengine ( )
```

7.23.3.17 get_states() [1/2]

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

Powerset of a gene's possible states.

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

Returns

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

Definition at line 665 of file defm-meat.hpp.

7.23.3.18 get_states() [2/2]

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

Powerset of a gene's possible states.

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

Returns

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

7.23.3.19 get_support_fun() [1/2]

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

Definition at line 661 of file defm-meat.hpp.

7.23.3.20 get_support_fun() [2/2]

```
phylocounters::PhyloSupport* Geese::get_support_fun ( )
```

7.23.3.21 inherit_support() [1/2]

Definition at line 230 of file defm-meat.hpp.

7.23.3.22 inherit_support() [2/2]

7.23.3.23 init() [1/2]

```
void Geese::init (
          unsigned int bar_width = BARRY_PROGRESS_BAR_WIDTH ) [inline]
```

Definition at line 103 of file defm-meat.hpp.

7.23.3.24 init() [2/2]

```
void Geese::init (
     unsigned int bar_width = BARRY_PROGRESS_BAR_WIDTH )
```

7.23.3.25 init_node() [1/2]

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

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

7.23.3.26 init_node() [2/2]

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

7.23.3.27 likelihood() [1/2]

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

7.23.3.28 likelihood() [2/2]

7.23.3.29 likelihood exhaust() [1/2]

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

7.23.3.30 likelihood_exhaust() [2/2]

7.23.3.31 nannotations() [1/2]

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

Number of annotations.

Definition at line 444 of file defm-meat.hpp.

7.23.3.32 nannotations() [2/2]

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

Number of annotations.

7.23.3.33 nfuns() [1/2]

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

Number of functions analyzed.

Definition at line 400 of file defm-meat.hpp.

7.23.3.34 nfuns() [2/2]

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

Number of functions analyzed.

7.23.3.35 nleafs() [1/2]

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

Number of leaf.

Definition at line 414 of file defm-meat.hpp.

7.23.3.36 nleafs() [2/2]

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

Number of leaf.

7.23.3.37 nnodes() [1/2]

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

Number of nodes (interior + leaf)

Definition at line 407 of file defm-meat.hpp.

7.23 Geese Class Reference 133

7.23.3.38 nnodes() [2/2]

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

Number of nodes (interior + leaf)

7.23.3.39 nterms() [1/2]

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

Number of terms included.

Definition at line 426 of file defm-meat.hpp.

7.23.3.40 nterms() [2/2]

```
unsigned int Geese::nterms ( ) const
```

Number of terms included.

7.23.3.41 observed_counts() [1/2]

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

Definition at line 495 of file defm-meat.hpp.

7.23.3.42 observed_counts() [2/2]

```
std::vector< std::vector<double> > Geese::observed_counts ( )
```

7.23.3.43 operator=() [1/4]

7.23.3.44 operator=() [2/4]

7.23.3.45 operator=() [3/4]

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

7.23.3.46 operator=() [4/4]

7.23.3.47 parse_polytomies() [1/2]

```
unsigned int Geese::parse_polytomies (
          bool verb = true,
          std::vector< size_t > * dist = nullptr ) const [inline], [noexcept]
```

Check polytomies and return the largest.

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

7.23.3.48 parse_polytomies() [2/2]

Check polytomies and return the largest.

7.23.3.49 predict() [1/2]

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

7.23.3.50 predict() [2/2]

7.23.3.51 predict_backend() [1/2]

< True if the array belongs to the set

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

7.23.3.52 predict_backend() [2/2]

7.23.3.53 predict_exhaust() [1/2]

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

7.23.3.54 predict_exhaust() [2/2]

7.23.3.55 predict_exhaust_backend() [1/2]

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

7.23.3.56 predict_exhaust_backend() [2/2]

7.23.3.57 predict_sim() [1/2]

```
std::vector< std::vector< double > > Geese::predict_sim (
    const std::vector< double > & par,
    bool only_annotated = false,
    unsigned int nsims = 10000u ) [inline]
```

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

7.23.3.58 predict_sim() [2/2]

7.23.3.59 print() [1/2]

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

Prints information about the DEFM.

Definition at line 629 of file defm-meat.hpp.

7.23 Geese Class Reference 137

7.23.3.60 print() [2/2]

```
void Geese::print ( ) const
```

Prints information about the GEESE.

7.23.3.61 print_observed_counts() [1/2]

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

Definition at line 566 of file defm-meat.hpp.

7.23.3.62 print_observed_counts() [2/2]

```
void Geese::print_observed_counts ( )
```

7.23.3.63 set_seed() [1/2]

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

7.23.3.64 set_seed() [2/2]

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

7.23.3.65 simulate() [1/2]

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

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

7.23.3.66 simulate() [2/2]

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

7.23.3.67 support_size() [1/2]

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

Number of unique sets of sufficient stats.

Definition at line 434 of file defm-meat.hpp.

7.23.3.68 support_size() [2/2]

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

Number of unique sets of sufficient stats.

7.23.3.69 update_annotations() [1/2]

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

Definition at line 258 of file defm-meat.hpp.

7.23.3.70 update_annotations() [2/2]

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

7.23.4 Member Data Documentation

7.23.4.1 delete_rengine

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

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

7.23.4.2 delete_support

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

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

7.23.4.3 initialized

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

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

7.23.4.4 map_to_nodes

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

Definition at line 94 of file defm-bones.hpp.

7.23.4.5 nfunctions

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

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

7.23.4.6 nodes

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

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

7.23.4.7 pset_loc

std::vector< std::vector< size_t > > > Geese::pset_loc

Locations of columns.

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

7.23.4.8 reduced_sequence

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

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

7.23.4.9 sequence

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

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

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

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

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

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

#include <model-bones.hpp>

Public Member Functions

- void set_rengine (std::mt19937 *rengine_, bool delete_=false)
- void set seed (unsigned int s)
- Model ()
- Model (uint size)
- Model (const Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > &Model ←
- Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > & operator= (const Model < Array Type, Data Counter Type, Data Rule Type, Data Rule Dyn Type > &Model)
- ∼Model ()
- void store_psets () noexcept
- void set_keygen (std::function < std::vector < double > (const Array_Type &) > keygen_)
- std::vector< double > gen_key (const Array_Type &Array_)
- uint add_array (const Array_Type &Array_, bool force_new=false)

Adds an array to the support of not already included.

- · void print stats (uint i) const
- · void print () const

Prints information about the model.

- Array_Type sample (const Array_Type &Array_, const std::vector< double > ¶ms={})
- Array_Type sample (const uint &i, const std::vector< double > ¶ms)
- double conditional_prob (const Array_Type &Array_, const std::vector< double > ¶ms, unsigned int i, unsigned int j)

Conditional probability ("Gibbs sampler")

- const std::mt19937 * get_rengine () const
- Counters < Array Type, Data Counter Type > * get counters ()
- Rules< Array_Type, Data_Rule_Type > * get_rules ()
- Rules< Array Type, Data Rule Dyn Type > * get rules dyn ()
- Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type > * get_support_fun ()

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_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)
- 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 fun type< Array Type, Data Rule Type > count fun , Data Rule Type data)
- 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_fun_type< Array_Type, Data_Rule_Dyn_Type > count_fun_, Data_Rule_Dyn
 — Type data_)
- void set rules dyn (Rules< Array Type, Data Rule Dyn Type > *rules)

Likelihood functions.

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

Parameters

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

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

Extract elements by index

Parameters

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

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

Size of the model

Number of different supports included in the model

This will return the size of stats_target.

Returns

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

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

Raw pointers to the support and target statistics.

- std::vector< std::vector< double > > * get stats support ()
- std::vector< unsigned int > * get_arrays2support ()
- std::vector< std::vector< Array_Type >> * get_pset_arrays ()
- std::vector< std::vector< double >>> * get_pset_stats ()

Statistics of the support(s)

- std::vector< std::vector< double > > * get_pset_probs ()
- void set_transform_model (std::function< std::vector< double >(double *, unsigned int)> fun, std::vector< std::string > names)

Set the transform_model_fun object.

• std::vector< double > transform_model (double *data, unsigned int k)

7.24.1 Detailed Description

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

class Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >

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

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

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

Template Parameters

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

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

7.24.2 Constructor & Destructor Documentation

7.24.2.1 Model() [1/3]

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

7.24.2.2 Model() [2/3]

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

7.24.2.3 Model() [3/3]

7.24.2.4 ~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 175 of file model-bones.hpp.

7.24.3 Member Function Documentation

7.24.3.1 add_array()

Adds an array to the support of not already included.

Parameters

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

Returns

The number of the array.

7.24.3.2 add_counter() [1/2]

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

7.24.3.3 add_counter() [2/2]

7.24.3.4 add_rule() [1/2]

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
void Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::add_rule (
Rule< Array_Type, Data_Rule_Type > & rule )
```

7.24.3.5 add_rule() [2/2]

7.24.3.6 add_rule_dyn() [1/2]

7.24.3.7 add_rule_dyn() [2/2]

7.24.3.8 colnames()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::string > Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_\times
Dyn_Type >::colnames ( ) const
```

7.24.3.9 conditional_prob()

Conditional probability ("Gibbs sampler")

Computes the conditional probability of observing $P\{Y(i,j) = | Y^{\wedge}C, \text{ theta}\}$, i.e., the probability of observing the entry Y(i,j) equal to one given the rest of the array.

Parameters

Array←	Array to check
_	
params	Vector of parameters
i	Row entry
j	Column entry

Returns

double The conditional probability

7.24.3.10 gen key()

7.24.3.11 get_arrays2support()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< unsigned int >* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_←
Rule_Dyn_Type >::get_arrays2support ()
```

7.24.3.12 get_counters()

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

7.24.3.13 get_norm_const()

7.24.3.14 get pset()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\leftarrow Rule_Type = bool, typename Data_Rule_Dyn_Type = bool> const std::vector< Array_Type >* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data\leftarrow _Rule_Dyn_Type >::get_pset ( const uint & i )
```

7.24.3.15 get_pset_arrays()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_\times
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
std::vector< std::vector< Array_Type > >* Model< Array_Type, Data_Counter_Type, Data_Rule_\times
Type, Data_Rule_Dyn_Type >::get_pset_arrays ()
```

7.24.3.16 get_pset_probs()

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

7.24.3.17 get_pset_stats() [1/2]

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

Statistics of the support(s)

7.24.3.18 get pset stats() [2/2]

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

7.24.3.19 get_rengine()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
const std::mt19937* Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type
>::get_rengine ( ) const
```

7.24.3.20 get_rules()

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

7.24.3.21 get_rules_dyn()

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

7.24.3.22 get_stats_support()

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

7.24.3.23 get_stats_target()

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

Raw pointers to the support and target statistics.

The support of the model is stored as a vector of vector < double>. Each element of it contains the support for an specific type of array included. It represents an array of size $(k + 1) \times n$ unique elements, with the data stored by-row. The last element of each entry corresponds to the weights, i.e., the frequency with which such sufficient statistics are observed in the support.

7.24.3.24 get support fun()

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

7.24.3.25 likelihood() [1/3]

7.24.3.26 likelihood() [2/3]

7.24.3.27 likelihood() [3/3]

7.24.3.28 likelihood_total()

7.24.3.29 nterms()

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

7.24.3.30 operator=()

7.24.3.31 print()

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

Prints information about the model.

7.24.3.32 print_stats()

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

7.24.3.33 sample() [1/2]

7.24.3.34 sample() [2/2]

7.24.3.35 set_counters()

7.24.3.36 set_keygen()

7.24.3.37 set_rengine()

Definition at line 145 of file model-bones.hpp.

7.24.3.38 set_rules()

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

7.24.3.39 set_rules_dyn()

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

7.24.3.40 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 155 of file model-bones.hpp.

7.24.3.41 set transform model()

Set the transform_model_fun object.

The transform model function is used to transform the data

Parameters

data	
target	
n_arrays	
arrays2support	

7.24.3.42 size()

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

7.24.3.43 size_unique()

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

7.24.3.44 store_psets()

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

7.24.3.45 support_size()

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

7.24.3.46 transform_model()

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

• include/barry/model-bones.hpp

7.25 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

7.25.1 Detailed Description

Data class used to store arbitrary uint or double vectors.

Definition at line 56 of file network.hpp.

7.25.2 Constructor & Destructor Documentation

7.25.2.1 NetCounterData() [1/2]

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

Definition at line 62 of file network.hpp.

7.25.2.2 NetCounterData() [2/2]

Definition at line 63 of file network.hpp.

7.25.2.3 ~NetCounterData()

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

Definition at line 68 of file network.hpp.

7.25.3 Member Data Documentation

7.25.3.1 indices

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

Definition at line 59 of file network.hpp.

7.25.3.2 numbers

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

Definition at line 60 of file network.hpp.

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

• include/barry/counters/network.hpp

7.26 NetworkData Class Reference

Data class for Networks.

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

Public Member Functions

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

Constructor using a single attribute.

- NetworkData (std::vector < std::vector < double > > vertex_attr_, bool directed_=true)
 Constructor using multiple attributes.
- ∼NetworkData ()

Public Attributes

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

7.26.1 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 19 of file network.hpp.

7.26.2 Constructor & Destructor Documentation

7.26.2.1 NetworkData() [1/3]

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

Definition at line 25 of file network.hpp.

7.26.2.2 NetworkData() [2/3]

Constructor using a single attribute.

Parameters

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

Definition at line 33 of file network.hpp.

7.26.2.3 NetworkData() [3/3]

Constructor using multiple attributes.

Parameters

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

Definition at line 45 of file network.hpp.

7.26.2.4 ~NetworkData()

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

Definition at line 51 of file network.hpp.

7.26.3 Member Data Documentation

7.26.3.1 directed

bool NetworkData::directed = true

Definition at line 22 of file network.hpp.

7.26.3.2 vertex_attr

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

Definition at line 23 of file network.hpp.

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

• include/barry/counters/network.hpp

7.27 Node Class Reference

A single node for the model.

#include <geese-node-bones.hpp>

Collaboration diagram for Node:



7.27 Node Class Reference 159

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.

7.27.1 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.

7.27.2 Constructor & Destructor Documentation

7.27.2.1 Node() [1/5]

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

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

7.27.2.2 Node() [2/5]

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

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

7.27.2.3 Node() [3/5]

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

7.27.2.4 Node() [4/5]

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

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

7.27.2.5 Node() [5/5]

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

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

7.27 Node Class Reference 161

7.27.2.6 ∼Node()

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

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

7.27.3 Member Function Documentation

7.27.3.1 get_parent()

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

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

7.27.3.2 is_leaf()

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

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

7.27.3.3 noffspring()

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

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

7.27.4 Member Data Documentation

7.27.4.1 annotations

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

Observed annotations (only defined for Geese)

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

7.27.4.2 array

```
phylocounters::PhyloArray Node::array
```

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

7.27.4.3 arrays

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

Arrays given all possible states.

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

7.27.4.4 duplication

bool Node::duplication

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

7.27.4.5 id

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

Id of the node (as specified in the input)

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

7.27.4.6 narray

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

ID of the array in the model.

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

7.27 Node Class Reference 163

7.27.4.7 offspring

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

Offspring nodes.

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

7.27.4.8 ord

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

Order in which the node was created.

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

7.27.4.9 parent

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

Parent node.

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

7.27.4.10 probability

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

The probability of observing each state.

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

7.27.4.11 subtree_prob

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

Induced subtree probabilities.

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

7.27.4.12 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

7.28 NodeData Class Reference

Data definition for the PhyloArray class.

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

Public Member Functions

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

Public Attributes

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

• bool duplication = true

7.28.1 Detailed Description

Data definition for the PhyloArray class.

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

This holds basic information about a given node.

Definition at line 38 of file phylo.hpp.

7.28.2 Constructor & Destructor Documentation

7.28.2.1 NodeData()

Definition at line 58 of file phylo.hpp.

7.28.3 Member Data Documentation

7.28.3.1 blengths

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

Branch length.

Definition at line 44 of file phylo.hpp.

7.28.3.2 duplication

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

Definition at line 54 of file phylo.hpp.

7.28.3.3 states

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

State of the parent node.

Definition at line 49 of file phylo.hpp.

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

• include/barry/counters/phylo.hpp

7.29 PhyloCounterData Class Reference

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

Public Member Functions

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

7.29.1 Detailed Description

Definition at line 69 of file phylo.hpp.

7.29.2 Constructor & Destructor Documentation

7.29.2.1 PhyloCounterData() [1/2]

Definition at line 75 of file phylo.hpp.

7.29.2.2 PhyloCounterData() [2/2]

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

Definition at line 80 of file phylo.hpp.

7.29.3 Member Function Documentation

7.29.3.1 at()

Definition at line 82 of file phylo.hpp.

7.29.3.2 begin()

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

Definition at line 90 of file phylo.hpp.

7.29.3.3 empty()

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

Definition at line 93 of file phylo.hpp.

7.29.3.4 end()

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

Definition at line 91 of file phylo.hpp.

7.29.3.5 get_counters()

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

Definition at line 94 of file phylo.hpp.

7.29.3.6 operator()()

Definition at line 83 of file phylo.hpp.

7.29.3.7 operator[]()

```
uint PhyloCounterData::operator[] (
          uint d) [inline]
```

Definition at line 84 of file phylo.hpp.

7.29.3.8 push_back()

Definition at line 86 of file phylo.hpp.

7.29.3.9 reserve()

Definition at line 85 of file phylo.hpp.

7.29.3.10 shrink_to_fit()

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

Definition at line 87 of file phylo.hpp.

7.29.3.11 size()

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

Definition at line 88 of file phylo.hpp.

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

• include/barry/counters/phylo.hpp

7.30 PhyloRuleDynData Class Reference

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

Public Member Functions

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

Public Attributes

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

7.30.1 Detailed Description

Definition at line 2147 of file phylo.hpp.

7.30.2 Constructor & Destructor Documentation

7.30.2.1 PhyloRuleDynData()

Definition at line 2154 of file phylo.hpp.

7.30.2.2 ~PhyloRuleDynData()

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

Definition at line 2163 of file phylo.hpp.

7.30.3 Member Data Documentation

7.30.3.1 counts

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

Definition at line 2149 of file phylo.hpp.

7.30.3.2 duplication

```
uint PhyloRuleDynData::duplication
```

Definition at line 2153 of file phylo.hpp.

7.30.3.3 lb

```
uint PhyloRuleDynData::lb
```

Definition at line 2151 of file phylo.hpp.

7.30.3.4 pos

uint PhyloRuleDynData::pos

Definition at line 2150 of file phylo.hpp.

7.30.3.5 ub

uint PhyloRuleDynData::ub

Definition at line 2152 of file phylo.hpp.

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

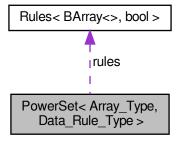
• include/barry/counters/phylo.hpp

7.31 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_fun_type< Array_Type, Data_Rule_Type > count_fun_, Data_Rule_Type data_)

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 < size_t > coordinates_free
- std::vector< size_t > coordinates_locked
- size_t n_free
- size_t n_locked

7.31.1 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.

7.31.2 Constructor & Destructor Documentation

7.31.2.1 PowerSet() [1/3]

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

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

7.31.2.2 PowerSet() [2/3]

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

7.31.2.3 PowerSet() [3/3]

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

7.31.2.4 ∼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.

7.31.3 Member Function Documentation

7.31.3.1 add_rule() [1/2]

Definition at line 175 of file powerset-meat.hpp.

7.31.3.2 add_rule() [2/2]

Definition at line 184 of file powerset-meat.hpp.

7.31.3.3 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 74 of file powerset-bones.hpp.

7.31.3.4 calc()

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

Definition at line 146 of file powerset-meat.hpp.

7.31.3.5 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 75 of file powerset-bones.hpp.

7.31.3.6 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 73 of file powerset-bones.hpp.

7.31.3.7 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 72 of file powerset-bones.hpp.

7.31.3.8 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.

7.31.3.9 operator[]()

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

7.31.3.10 reset()

Definition at line 162 of file powerset-meat.hpp.

7.31.3.11 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 76 of file powerset-bones.hpp.

7.31.4 Member Data Documentation

7.31.4.1 coordinates_free

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

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

7.31.4.2 coordinates locked

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

Definition at line 33 of file powerset-bones.hpp.

7.31.4.3 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 25 of file powerset-bones.hpp.

7.31.4.4 EmptyArray

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

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

7.31.4.5 M

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

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

7.31.4.6 N

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

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

7.31.4.7 n_free

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

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

7.31.4.8 n locked

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

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

7.31.4.9 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 26 of file powerset-bones.hpp.

7.31.4.10 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 29 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

7.32 Progress Class Reference

A simple progress bar.

```
#include progress.hpp>
```

Public Member Functions

```
• Progress (int n_, int width_)
```

- ∼Progress ()
- void next ()
- void end ()

7.32.1 Detailed Description

A simple progress bar.

Definition at line 11 of file progress.hpp.

7.32.2 Constructor & Destructor Documentation

7.32.2.1 Progress()

```
Progress::Progress (
          int n_,
          int width_ ) [inline]
```

Definition at line 30 of file progress.hpp.

7.32.2.2 ∼Progress()

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

Definition at line 23 of file progress.hpp.

7.32.3 Member Function Documentation

7.32.3.1 end()

```
void Progress::end ( ) [inline]
```

Definition at line 52 of file progress.hpp.

7.32.3.2 next()

```
void Progress::next ( ) [inline]
```

Definition at line 41 of file progress.hpp.

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

• include/barry/progress.hpp

7.33 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_← dat_	When true, the Rule destructor will delete the pointer, if defined.

- Rule ()
- Rule (Rule_fun_type< Array_Type, Data_Type > fun_, Data_Type dat_)

7.33.1 Detailed Description

```
\label{template} $$ \ensuremath{\sf template}$$ < typename \ensuremath{\sf Array\_Type}$ = BArray<>, typename \ensuremath{\sf Data\_Type}$ = bool> class \ensuremath{\sf Rule}< Array\_Type, \ensuremath{\sf 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 22 of file rules-bones.hpp.

7.33.2 Constructor & Destructor Documentation

7.33.2.1 Rule() [1/2]

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

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

7.33.2.2 Rule() [2/2]

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

7.33.2.3 ∼Rule()

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

Definition at line 47 of file rules-bones.hpp.

7.33.3 Member Function Documentation

7.33.3.1 D()

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

Read/Write access to the data.

7.33.3.2 operator()()

Definition at line 42 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

7.34 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 get_seq (const Array_Type &a, std::vector< size_t > *free, std::vector< size_t > *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_fun_type< Array_Type, Data_Type > rule_, Data_Type data_)

7.34.1 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 62 of file rules-bones.hpp.

7.34.2 Constructor & Destructor Documentation

7.34.2.1 Rules() [1/2]

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

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

7.34.2.2 Rules() [2/2]

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

7.34.2.3 ∼Rules()

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

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

7.34.3 Member Function Documentation

7.34.3.1 add_rule() [1/2]

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

7.34.3.2 add_rule() [2/2]

Definition at line 57 of file rules-meat.hpp.

7.34.3.3 get_seq()

Computes the sequence of free and locked cells in an ${\ensuremath{\mathsf{BArray}}}.$

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 88 of file rules-meat.hpp.

7.34.3.4 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 72 of file rules-meat.hpp.

7.34.3.5 operator=()

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

7.34.3.6 size()

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

Definition at line 75 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

7.35 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 (const StatsCounter< Array_Type, Data_Type > &counter)

Copy constructor.

· 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 ()
- std::vector< std::string > get_names () const
- std::vector< std::string > get_descriptions () const
- size_t size () const

7.35.1 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 19 of file statscounter-bones.hpp.

7.35.2 Constructor & Destructor Documentation

7.35.2.1 StatsCounter() [1/3]

Creator of a StatsCounter

Parameters

Array⇔	A const pointer to a BArray.
_	

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

7.35.2.2 StatsCounter() [2/3]

Copy constructor.

Parameters

counter

7.35.2.3 StatsCounter() [3/3]

```
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 64 of file statscounter-bones.hpp.

7.35.2.4 ~StatsCounter()

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

7.35.3 Member Function Documentation

7.35.3.1 add_counter() [1/2]

7.35.3.2 add_counter() [2/2]

7.35.3.3 count_all()

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

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

7.35.3.4 count_current()

7.35.3.5 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.

7.35.3.6 get counters()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
Counters<Array_Type,Data_Type>* StatsCounter< Array_Type, Data_Type >::get_counters ( )
```

7.35.3.7 get_descriptions()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
std::vector< std::string > StatsCounter< Array_Type, Data_Type >::get_descriptions ( ) const
```

7.35.3.8 get_names()

```
template<typename Array_Type = BArray<>, typename Data_Type = bool>
std::vector< std::string > StatsCounter< Array_Type, Data_Type >::get_names ( ) const
```

7.35.3.9 reset array()

Changes the reference array for the counting.

Parameters

Array⇔	A pointer to an array of class Array_Type.

7.35.3.10 set_counters()

7.35.3.11 size()

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

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

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

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

7.36 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.

- std::vector< double > get_counts () 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_)
- bool eval_rules_dyn (const std::vector< double > &counts, const uint &i, const uint &j)

Public Attributes

- uint N
- uint M
- bool delete_counters = true
- bool delete rules = true
- bool delete rules dyn = true
- uint max num elements = BARRY MAX NUM ELEMENTS
- std::vector< double > current stats
- std::vector< size t > coordinates free
- std::vector< size t > coordinates locked
- · size t coordiantes n free
- · size t coordiantes n locked
- std::vector< double > change_stats
- std::vector< size_t > hashes
- std::vector< bool > hashes_initialized
- size_t n_counters

7.36.1 Detailed Description

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

 ${\it class \ Support} {\it <Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type} >$

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.

7.36.2 Constructor & Destructor Documentation

7.36.2.1 Support() [1/3]

Constructor passing a reference Array.

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

7.36.2.2 Support() [2/3]

Constructor specifying the dimensions of the array (empty).

Definition at line 89 of file support-bones.hpp.

7.36.2.3 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 96 of file support-bones.hpp.

7.36.2.4 ~Support()

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

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

7.36.3 Member Function Documentation

7.36.3.1 add_counter() [1/2]

7.36.3.2 add_counter() [2/2]

7.36.3.3 add_rule() [1/2]

7.36.3.4 add rule() [2/2]

7.36.3.5 add_rule_dyn() [1/2]

7.36.3.6 add_rule_dyn() [2/2]

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

7.36.3.7 calc()

```
template<typename Array_Type = BArray<>, typename Data_Counter_Type = bool, typename Data_←
Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>

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

Computes the entire support.

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

Parameters

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

7.36.3.8 eval rules dyn()

7.36.3.9 get_counters()

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

Vector of couter functions.

7.36.3.10 get counts()

```
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 >::get_counts ( ) const
```

7.36.3.11 get current stats()

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

List current statistics.

7.36.3.12 get_data()

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

7.36.3.13 get_rules()

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

Vector of static rules (cells to iterate).

7.36.3.14 get_rules_dyn()

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

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

7.36.3.15 init_support()

7.36.3.16 print()

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

7.36.3.17 reset_array() [1/2]

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

7.36.3.18 reset_array() [2/2]

7.36.3.19 set_counters()

7.36.3.20 set rules()

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

7.36.3.21 set_rules_dyn()

7.36.4 Member Data Documentation

7.36.4.1 change 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 >::change_stats
```

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

7.36.4.2 coordiantes_n_free

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

Rule_Type = bool, typename Data_Rule_Dyn_Type = bool>
size_t Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >::coordiantes\( \cdot\)
_n_free
```

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

7.36.4.3 coordiantes_n_locked

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

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

7.36.4.4 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< size_t > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn←
_Type >::coordinates_free
```

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

7.36.4.5 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< size_t > Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn←
_Type >::coordinates_locked
```

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

7.36.4.6 current stats

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

Definition at line 68 of file support-bones.hpp.

7.36.4.7 delete_counters

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

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

7.36.4.8 delete_rules

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

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

7.36.4.9 delete_rules_dyn

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

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

7.36.4.10 hashes

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

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

7.36.4.11 hashes_initialized

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

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

7.36.4.12 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 61 of file support-bones.hpp.

7.36.4.13 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 65 of file support-bones.hpp.

7.36.4.14 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 61 of file support-bones.hpp.

7.36.4.15 n_counters

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

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

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

• include/barry/support-bones.hpp

7.37 vecHasher < T > Struct Template Reference

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

Public Member Functions

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

7.37.1 Detailed Description

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

Definition at line 106 of file typedefs.hpp.

7.37.2 Member Function Documentation

7.37.2.1 operator()()

Definition at line 109 of file typedefs.hpp.

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

include/barry/typedefs.hpp

Chapter 8

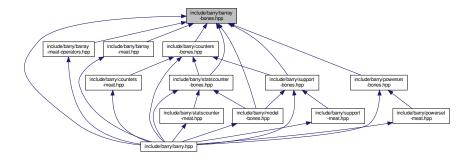
File Documentation

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

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



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



Classes

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

Macros

• #define BARRAY_BONES_HPP 1

8.1.1 Macro Definition Documentation

8.1.1.1 BARRAY_BONES_HPP

#define BARRAY_BONES_HPP 1

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

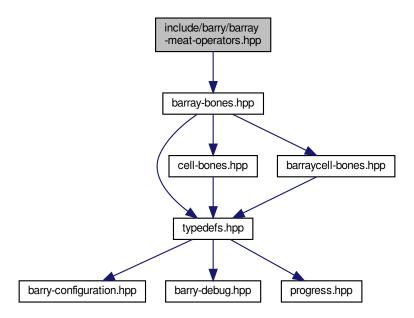
8.2 include/barry/barray-iterator.hpp File Reference

Classes

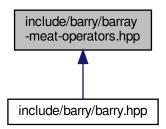
 $\bullet \ \, {\sf class\ ConstBArrayRowlter} < {\sf Cell_Type}, \, {\sf Data_Type} >$

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 BARRAY_TYPE() BArray<Cell_Type, Data_Type>
- #define BARRAY_TEMPLATE_ARGS() < typename Cell_Type, typename Data_Type>
- #define BARRAY_TEMPLATE(a, b) template BARRAY_TEMPLATE_ARGS() inline a BARRAY_TYPE()::b
- #define ROW(a) this->el_ij[a]
- #define COL(a) this->el_ji[a]

Functions

- template BARRAY_TEMPLATE_ARGS () inline void checkdim_(const BARRAY_TYPE() &lhs
- template const BARRAY_TYPE () &rhs)
- BARRAY_TEMPLATE (BARRAY_TYPE()&, operator+=)(const BArray< Cell_Type
- for (uint i=0u;i< nrow();++i) for(uint j=0u = el[POS(i, j)]
- j< ncol();++j) this-> operator() (i, j)+
- BARRAY_TEMPLATE (BARRAY_TYPE()&, operator+=)(const Cell_Type &rhs)
- BARRAY_TEMPLATE (BARRAY_TYPE()&, operator-=)(const BArray< Cell_Type
- BARRAY_TEMPLATE (BARRAY_TYPE()&, operator-=)(const Cell_Type &rhs)
- BARRAY TEMPLATE (BARRAY TYPE()&, operator*=)(const Cell Type &rhs)
- BARRAY_TEMPLATE (BARRAY_TYPE()&, operator/=)(const Cell_Type &rhs)

Variables

- Data_Type & rhs
- return * this

8.3.1 Macro Definition Documentation

8.3.1.1 BARRAY_TEMPLATE

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

8.3.1.2 BARRAY_TEMPLATE_ARGS

```
template BARRAY_TEMPLATE_ARGS() <typename Cell_Type, typename Data_Type>
```

Definition at line 9 of file barray-meat-operators.hpp.

8.3.1.3 BARRAY_TYPE

```
template Data_Type BARRAY_TYPE( ) BArray<Cell_Type, Data_Type>
```

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

8.3.1.4 BARRY_BARRAY_MEAT_OPERATORS_HPP

```
#define BARRY_BARRAY_MEAT_OPERATORS_HPP 1
```

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

8.3.1.5 COL

Definition at line 15 of file barray-meat-operators.hpp.

8.3.1.6 ROW

Definition at line 14 of file barray-meat-operators.hpp.

8.3.2 Function Documentation

8.3.2.1 BARRAY_TEMPLATE() [1/6]

Definition at line 88 of file barray-meat-operators.hpp.

8.3.2.2 BARRAY_TEMPLATE() [2/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator+ ) const
```

8.3.2.3 BARRAY_TEMPLATE() [3/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator+ ) const &
```

Definition at line 46 of file barray-meat-operators.hpp.

8.3.2.4 BARRAY_TEMPLATE() [4/6]

8.3.2.5 BARRAY_TEMPLATE() [5/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator- ) const &
```

Definition at line 75 of file barray-meat-operators.hpp.

8.3.2.6 BARRAY_TEMPLATE() [6/6]

```
BARRAY_TEMPLATE (
          BARRAY_TYPE()& ,
          operator/ ) const &
```

Definition at line 105 of file barray-meat-operators.hpp.

8.3.2.7 BARRAY_TEMPLATE_ARGS()

```
template BARRAY_TEMPLATE_ARGS ( ) const &
```

8.3.2.8 BARRAY_TYPE()

```
template const BARRAY_TYPE ( ) &
```

Definition at line 20 of file barray-meat-operators.hpp.

8.3.2.9 for()

```
for ( ) = el[POS(i, j)] [pure virtual]
```

Definition at line 66 of file barray-meat-operators.hpp.

8.3.2.10 operator()()

8.3.3 Variable Documentation

8.3.3.1 rhs

```
Data_Type & rhs
Initial value:
{
    checkdim_(*this, rhs)
```

Definition at line 33 of file barray-meat-operators.hpp.

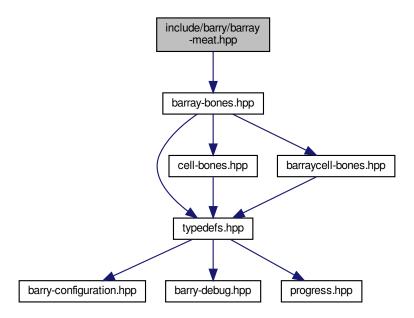
8.3.3.2 this

```
return * this
```

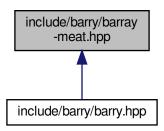
Definition at line 43 of file barray-meat-operators.hpp.

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 BARRAY_TYPE() BArray<Cell_Type, Data_Type>
- #define BARRAY_TEMPLATE_ARGS() < typename Cell_Type, typename Data_Type>
- #define BARRAY_TEMPLATE(a, b) template BARRAY_TEMPLATE_ARGS() inline a BARRAY_TYPE()::b
- #define ROW(a) this->el_ij[a]
- #define COL(a) this->el_ji[a]

Functions

```
• BARRAY TEMPLATE (, BArray)(uint N_

 el_ij resize (N)

• el_ji resize (M)
• for (uint i=0u;i < source.size();++i)

    Data Type bool M (Array .M)

    BARRAY_TEMPLATE (BARRAY_TYPE() &, operator=)(const BArray< Cell_Type</li>

    BARRAY_TEMPLATE (, BArray)(BARRAY_TYPE() &&x) noexcept

    BARRAY TEMPLATE (BARRAY TYPE() &, operator=)(BARRAY TYPE() &&x) noexcept

• BARRAY TEMPLATE (bool, operator==)(const BARRAY TYPE() & Array )

    BARRAY_TEMPLATE (,~BArray)()

    BARRAY_TEMPLATE (void, set_data)(Data_Type *data_

    BARRAY TEMPLATE (Data Type *, D)()

• BARRAY_TEMPLATE (void, out_of_range)(uint i

    BARRAY_TEMPLATE (Cell_Type, get_cell)(uint i

    if (ROW(i).size()==0u) return(Cell Type) 0.0

• if (search !=ROW(i).end()) return search -> second.value
• return (Cell_Type) 0.0

    BARRAY_TEMPLATE (std::vector< Cell_Type >, get_row_vec)(uint i

    std::vector< Cell Type > ans (ncol(),(Cell Type) false)

    for (const auto &iter :row(i, false)) ans[iter.first]

    BARRAY_TEMPLATE (void, get_row_vec)(std

    BARRAY_TEMPLATE (BARRAY_TYPE() &, operator-=)(const std

    BARRAY_TEMPLATE (void, insert_cell)(uint i

• if (check_exists)

    COL (j).emplace(i

• & ROW (i)[j])

    BARRAY_TEMPLATE (void, swap_cells)(uint i0

if (report !=nullptr)(*report)
• if (check0 &check1)
• else if (!check0 &check1)

    else if (check0 &!check1)

    BARRAY_TEMPLATE (void, toggle_cell)(uint i

    BARRAY TEMPLATE (void, swap rows)(uint i0

• if (ROW(i0).size()==0u) move0
if (ROW(i1).size()==0u) move1
• if (!move0 &&!move1) return

    ROW (i0).swap(ROW(i1))

    BARRAY TEMPLATE (void, swap cols)(uint j0

    if (COL(j0).size()==0u) check0

if (COL(j1).size()==0u) check1

    if (check0 &&check1)

• else if (check0 &&!check1)

    else if (!check0 &&check1)

    BARRAY_TEMPLATE (void, zero_row)(uint i

for (auto row=row0.begin();row !=row0.end();++row) rm_cell(i

    BARRAY TEMPLATE (void, zero col)(uint i

• if (COL(j).size()==0u) return

    BARRAY TEMPLATE (void, transpose)()

    BARRAY_TEMPLATE (void, clear)(bool hard)

    BARRAY_TEMPLATE (void, resize)(uint N_

• if (M_< M) for(uint j = N_
```

Variables

```
    uint M
```

- uint const std::vector< uint > & source
- uint const std::vector< uint > const std::vector< uint > & target
- uint const std::vector< uint > const std::vector< cell_Type > & value
- uint const std::vector< uint > const std::vector< Cell Type > bool add
- if(source.size() !=value.size()) throw std N = N_
- M = M
- return
- Data_Type & Array_
- Data_Type bool copy_data
- bool delete_data_
- data = data
- delete_data = delete_data_
- · uint j const
- uint j
- auto search = ROW(i).find(j)
- return ans
- uint const Cell
 Cell_Type > & v
- uint const Cell< Cell_Type > bool check_bounds
- uint const Cell
 Cell_Type > bool bool check_exists
- else
- NCells
- uint j0
- uint uint i1
- uint uint uint j1
- uint uint bool int int * report
- auto row0 = ROW(i)
- row first
- · row false
- auto col0 = COL(j)

8.4.1 Macro Definition Documentation

8.4.1.1 BARRAY_TEMPLATE

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

8.4.1.2 BARRAY_TEMPLATE_ARGS

```
#define BARRAY_TEMPLATE_ARGS( ) <typename Cell_Type, typename Data_Type>
```

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

8.4.1.3 BARRAY_TYPE

```
#define BARRAY_TYPE( ) BArray<Cell_Type, Data_Type>
```

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

8.4.1.4 COL

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

8.4.1.5 ROW

Definition at line 20 of file barray-meat.hpp.

8.4.2 Function Documentation

8.4.2.1 ans()

8.4.2.2 BARRAY_TEMPLATE() [1/23]

```
BARRAY_TEMPLATE (

BArray ) && [noexcept]
```

Definition at line 230 of file barray-meat.hpp.

8.4.2.3 BARRAY_TEMPLATE() [2/23]

```
BARRAY_TEMPLATE (
BArray )
```

8.4.2.4 BARRAY_TEMPLATE() [3/23]

```
BARRAY_TEMPLATE ( \sim \textit{BArray} \ )
```

Definition at line 339 of file barray-meat.hpp.

8.4.2.5 BARRAY_TEMPLATE() [4/23]

Definition at line 586 of file barray-meat.hpp.

8.4.2.6 BARRAY_TEMPLATE() [5/23]

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

8.4.2.7 BARRAY_TEMPLATE() [6/23]

8.4.2.8 BARRAY_TEMPLATE() [7/23]

```
BARRAY_TEMPLATE (
          bool ,
          operator = = ) const &
```

Definition at line 321 of file barray-meat.hpp.

8.4.2.9 BARRAY_TEMPLATE() [8/23]

8.4.2.10 BARRAY_TEMPLATE() [9/23]

Definition at line 361 of file barray-meat.hpp.

8.4.2.11 BARRAY_TEMPLATE() [10/23]

```
BARRAY_TEMPLATE (
          std::vector< Cell_Type > ,
          get_row_vec )
```

8.4.2.12 BARRAY_TEMPLATE() [11/23]

```
BARRAY_TEMPLATE (
     void ,
     clear )
```

Definition at line 1119 of file barray-meat.hpp.

8.4.2.13 BARRAY_TEMPLATE() [12/23]

Definition at line 441 of file barray-meat.hpp.

8.4.2.14 BARRAY_TEMPLATE() [13/23]

8.4.2.15 BARRAY_TEMPLATE() [14/23]

8.4.2.16 BARRAY_TEMPLATE() [15/23]

```
BARRAY_TEMPLATE (
            void ,
            resize )
```

8.4.2.17 BARRAY_TEMPLATE() [16/23]

8.4.2.18 BARRAY_TEMPLATE() [17/23]

8.4.2.19 BARRAY_TEMPLATE() [18/23]

8.4.2.20 BARRAY_TEMPLATE() [19/23]

```
BARRAY_TEMPLATE (
     void ,
     swap_rows )
```

8.4.2.21 BARRAY_TEMPLATE() [20/23]

8.4.2.22 BARRAY_TEMPLATE() [21/23]

Definition at line 1058 of file barray-meat.hpp.

8.4.2.23 BARRAY_TEMPLATE() [22/23]

8.4.2.24 BARRAY_TEMPLATE() [23/23]

8.4.2.25 COL()

```
COL (
```

```
8.4.2.26 for() [1/3]
```

```
for (
     auto row = row0.begin();row !=row0.end();++row )
```

8.4.2.27 for() [2/3]

8.4.2.28 for() [3/3]

```
for ( )
```

Definition at line 51 of file barray-meat.hpp.

8.4.2.29 if() [1/17]

```
else if (
     !check0 && check1 )
```

Definition at line 997 of file barray-meat.hpp.

8.4.2.30 if() [2/17]

```
else if (
    !check0 & check1 )
```

Definition at line 845 of file barray-meat.hpp.

8.4.2.31 if() [3/17]

```
if (
    !move0 &&! move1 )
```

8.4.2.32 if() [4/17]

Definition at line 853 of file barray-meat.hpp.

```
8.4.2.33 if() [5/17]
```

```
else if ( check0 &&! check1)
```

Definition at line 988 of file barray-meat.hpp.

```
8.4.2.34 if() [6/17]
```

```
if ( check0 && check1)
```

Definition at line 961 of file barray-meat.hpp.

8.4.2.35 if() [7/17]

```
if ( {\tt check0~\&~\it check1~)}
```

Definition at line 827 of file barray-meat.hpp.

8.4.2.36 if() [8/17]

```
else if (
          check_exists = = CHECK::BOTH )
```

Definition at line 668 of file barray-meat.hpp.

8.4.2.37 if() [9/17]

```
if ( \label{eq:col} \mathtt{COL}(\mathtt{j}).\mathtt{size}(\mathtt{)} \ = = 0u \ \mathtt{)}
```

```
8.4.2.38 if() [10/17]
```

```
if ( COL(j0).size() = =0u )
```

8.4.2.39 if() [11/17]

```
if ( \label{eq:col} \mathtt{COL}(\mathtt{j1}).\mathtt{size}() \ = = 0u \ )
```

8.4.2.40 if() [12/17]

```
else if ( ) = N_
```

Definition at line 86 of file barray-meat.hpp.

8.4.2.41 if() [13/17]

```
if (
    report ! = nullptr )
```

8.4.2.42 if() [14/17]

```
if ( \label{eq:row_row_row} \mbox{ROW(i).size()} \ = \ = 0 \mbox{$u$} \ )
```

8.4.2.43 if() [15/17]

```
if ( \label{eq:row_row} \mbox{ROW(i0).size()} \ = \ = \mbox{$0$$u$} \ )
```

8.4.2.44 if() [16/17]

```
if ( \label{eq:row_row_row} \text{ROW(i1).size()} \quad = = 0u \text{ )}
```

```
8.4.2.45 if() [17/17]
```

```
if (
    search ! = ROW(i).end() ) -> second.value
```

8.4.2.46 M()

```
Data_Type bool M ( \label{eq:Array_.} \text{Array}\_. \quad \textit{M} \ )
```

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

8.4.2.47 resize() [1/2]

```
el_ji resize (
M )
```

8.4.2.48 resize() [2/2]

```
el_ij resize (
N )
```

8.4.2.49 return()

8.4.2.50 ROW() [1/2]

```
& ROW ( i )
```

8.4.2.51 ROW() [2/2]

```
ROW ( i0 )
```

8.4.3 Variable Documentation

8.4.3.1 add

```
uint const std::vector< uint > const std::vector< uint > bool add

Initial value:
{
    if (source.size() != target.size())
        throw std::length_error("-source- and -target- don't match on length.")
```

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

8.4.3.2 ans

```
return ans
```

Definition at line 438 of file barray-meat.hpp.

8.4.3.3 Array_

```
Data_Type & Array_
```

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

8.4.3.4 check_bounds

```
Initial value:
{
    if (check_bounds) {
       out_of_range(i0,0u);
       out_of_range(i1,0u);
    }
    bool move0=true, move1=true
```

Definition at line 661 of file barray-meat.hpp.

8.4.3.5 check_exists

```
uint bool int check_exists
Initial value:
{
    if (check_bounds)
        out_of_range(i,j)
```

Definition at line 662 of file barray-meat.hpp.

8.4.3.6 col0

```
auto col0 = COL(j)
```

Definition at line 1050 of file barray-meat.hpp.

8.4.3.7 const

```
uint bool check_bounds const

Initial value:
{
    if (i >= N)
        throw std::range_error("The row is out of range.")
```

Definition at line 391 of file barray-meat.hpp.

8.4.3.8 copy_data

```
Data_Type bool copy_data
```

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

8.4.3.9 data

```
data = data_
```

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

8.4.3.10 delete_data

```
delete_data = delete_data_
```

Definition at line 355 of file barray-meat.hpp.

8.4.3.11 delete_data_

Definition at line 348 of file barray-meat.hpp.

8.4.3.12 else

Definition at line 692 of file barray-meat.hpp.

8.4.3.13 false

row false

Definition at line 1031 of file barray-meat.hpp.

8.4.3.14 first

row first

Definition at line 1031 of file barray-meat.hpp.

8.4.3.15 i1

```
uint i1
```

Definition at line 765 of file barray-meat.hpp.

8.4.3.16 j

```
uint j
```

Initial value:

```
if (init_fun == nullptr)
    return 0.0
```

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

8.4.3.17 j0

```
uint j0
```

Definition at line 764 of file barray-meat.hpp.

8.4.3.18 j1

```
uint j1
```

Definition at line 765 of file barray-meat.hpp.

8.4.3.19 M

```
M = M_{\underline{}}
```

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

8.4.3.20 M_

```
uint M_
Initial value:
{

    if (N_ < N)
        for (uint i = N_; i < N; ++i)
            zero_row(i, false)</pre>
```

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

8.4.3.21 N

```
if (source.size() != target.size()) throw std if (source.size() != value.size()) throw std N=N
```

Definition at line 43 of file barray-meat.hpp.

8.4.3.22 NCells

NCells

Definition at line 696 of file barray-meat.hpp.

8.4.3.23 report

```
uint uint uint bool int int* report
```

Definition at line 768 of file barray-meat.hpp.

8.4.3.24 return

return

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

8.4.3.25 row0

```
auto row0 = ROW(i)
```

Definition at line 1029 of file barray-meat.hpp.

8.4.3.26 search

```
auto search = ROW(i).find(j)
```

Definition at line 415 of file barray-meat.hpp.

8.4.3.27 source

```
uint const std::vector< uint > & source
```

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

8.4.3.28 target

```
uint const std::vector< uint > const std::vector< uint > & target
```

Definition at line 32 of file barray-meat.hpp.

8.4.3.29 v

```
uint Cell_Type v
```

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

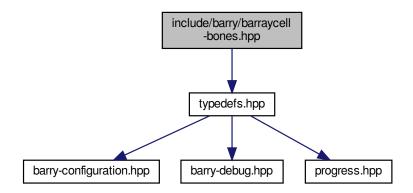
8.4.3.30 value

uint const std::vector< uint > const std::vector< uint > const std::vector< Cell_Type > & value

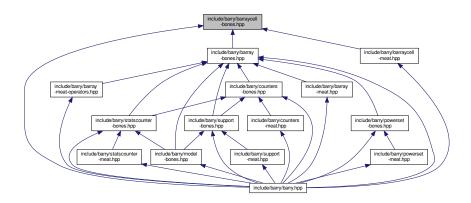
Definition at line 33 of file barray-meat.hpp.

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:



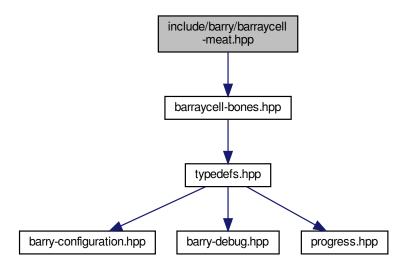
Classes

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

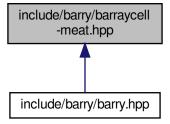
8.6 include/barry/barraycell-meat.hpp File Reference

#include "barraycell-bones.hpp"

Include dependency graph for barraycell-meat.hpp:



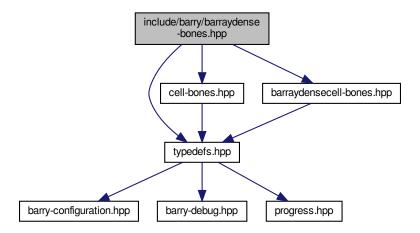
This graph shows which files directly or indirectly include this file:



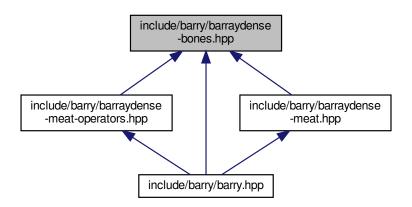
8.7 include/barry/barraydense-bones.hpp File Reference

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

Macros

• #define BARRY_BARRAYDENSE_BONES_HPP 1

8.7.1 Macro Definition Documentation

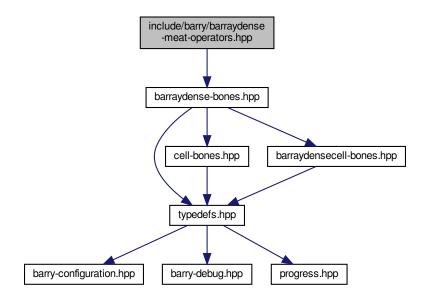
8.7.1.1 BARRY_BARRAYDENSE_BONES_HPP

#define BARRY_BARRAYDENSE_BONES_HPP 1

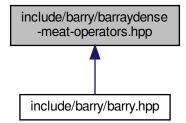
Definition at line 8 of file barraydense-bones.hpp.

8.8 include/barry/barraydense-meat-operators.hpp File Reference

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



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



Macros

- #define BARRY_BARRAYDENSE_MEAT_OPERATORS_HPP 1
- #define BDENSE_TYPE() BArrayDense<Cell_Type, Data_Type>
- #define BDENSE TEMPLATE ARGS() < typename Cell Type, typename Data Type>
- #define BDENSE TEMPLATE(a, b) template BDENSE TEMPLATE ARGS() inline a BDENSE TYPE()::b
- #define ROW(a) this->el ij[a]
- #define COL(a) this->el_ji[a]
- #define POS(a, b) (b)*N + (a)
- #define POS_N(a, b, c) (b)*(c) + (a)

Functions

- template BDENSE_TEMPLATE_ARGS () inline void checkdim_(const BDENSE_TYPE() &lhs
- template const BDENSE_TYPE () &rhs)
- BDENSE TEMPLATE (BDENSE TYPE()&, operator+=)(const BDENSE TYPE() &rhs)
- BDENSE TEMPLATE (BDENSE TYPE()&, operator-=)(const BDENSE TYPE() &rhs)
- BDENSE_TEMPLATE (BDENSE_TYPE()&, operator*=)(const Cell_Type &rhs)
- BDENSE_TEMPLATE (BDENSE_TYPE()&, operator/=)(const Cell_Type &rhs)

8.8.1 Macro Definition Documentation

8.8.1.1 BARRY_BARRAYDENSE_MEAT_OPERATORS_HPP

```
#define BARRY_BARRAYDENSE_MEAT_OPERATORS_HPP 1
```

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

8.8.1.2 BDENSE_TEMPLATE

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

8.8.1.3 BDENSE_TEMPLATE_ARGS

```
template BDENSE_TEMPLATE_ARGS() <typename Cell_Type, typename Data_Type>
```

Definition at line 9 of file barraydense-meat-operators.hpp.

8.8.1.4 BDENSE_TYPE

```
template Data_Type BDENSE_TYPE( ) BArrayDense<Cell_Type, Data_Type>
```

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

8.8.1.5 COL

Definition at line 15 of file barraydense-meat-operators.hpp.

8.8.1.6 POS

```
#define POS( \label{eq:a_b} a, \\ b \ ) \ (b)*N + (a)
```

Definition at line 16 of file barraydense-meat-operators.hpp.

8.8.1.7 POS N

Definition at line 17 of file barraydense-meat-operators.hpp.

8.8.1.8 ROW

Definition at line 14 of file barraydense-meat-operators.hpp.

8.8.2 Function Documentation

8.8.2.1 BDENSE_TEMPLATE() [1/4]

Definition at line 90 of file barraydense-meat-operators.hpp.

8.8.2.2 BDENSE_TEMPLATE() [2/4]

Definition at line 34 of file barraydense-meat-operators.hpp.

8.8.2.3 BDENSE_TEMPLATE() [3/4]

Definition at line 61 of file barraydense-meat-operators.hpp.

8.8.2.4 BDENSE_TEMPLATE() [4/4]

Definition at line 101 of file barraydense-meat-operators.hpp.

8.8.2.5 BDENSE_TEMPLATE_ARGS()

```
template BDENSE_TEMPLATE_ARGS ( ) const &
```

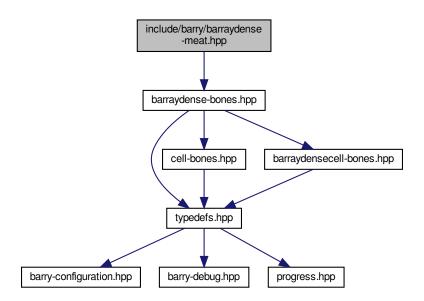
8.8.2.6 BDENSE_TYPE()

```
template const BDENSE_TYPE ( ) &
```

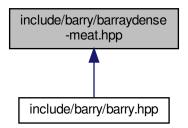
Definition at line 22 of file barraydense-meat-operators.hpp.

8.9 include/barry/barraydense-meat.hpp File Reference

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



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



Macros

- #define BDENSE_TYPE() BArrayDense<Cell_Type, Data_Type>
- #define BDENSE_TEMPLATE_ARGS() < typename Cell_Type, typename Data_Type>
- #define BDENSE_TEMPLATE(a, b) template BDENSE_TEMPLATE_ARGS() inline a BDENSE_TYPE()::b
- #define ROW(a) this->el_ij[a]
- #define COL(a) this->el_ji[a]
- #define POS(a, b) (b)*N + (a)
- #define POS_N(a, b, c) (b)*(c) + (a)
- #define ZERO_CELL static_cast<Cell_Type>(0.0)

Functions

```
    BDENSE_TEMPLATE (, BArrayDense)(uint N_

    el resize (N *M, ZERO_CELL)

• el_rowsums resize (N, ZERO_CELL)
• el_colsums resize (M, ZERO_CELL)

    for (uint i=0u;i< source.size();++i)</li>

    BDENSE_TEMPLATE (, BArrayDense)(const BDENSE_TYPE() &Array_

    bool M (Array .M)

• BDENSE_TEMPLATE (BDENSE_TYPE() &, operator=)(const BDENSE_TYPE() &Array_)
• BDENSE TEMPLATE (, BArrayDense)(BDENSE TYPE() &&x) noexcept

    BDENSE TEMPLATE (BDENSE TYPE() &, operator=)(BDENSE TYPE() &&x) noexcept

• BDENSE_TEMPLATE (bool, operator==)(const BDENSE_TYPE() &Array_)

    BDENSE TEMPLATE (, ~BArrayDense)()

    BDENSE_TEMPLATE (void, set_data)(Data_Type *data_

• BDENSE_TEMPLATE (Data_Type *, D)()
• BDENSE TEMPLATE (const Data Type *, D)() const
• BDENSE TEMPLATE (void, out of range)(uint i

    BDENSE TEMPLATE (Cell Type, get cell)(uint i

• BDENSE_TEMPLATE (std::vector< Cell_Type >, get_row_vec)(uint i

    std::vector< Cell Type > ans (ncol(), static cast< Cell Type >(false))

• BDENSE_TEMPLATE (void, get_row_vec)(std
• BDENSE TEMPLATE (Entries < Cell Type >, get entries)() const
• BDENSE TEMPLATE (bool, is empty)(uint i

    BDENSE TEMPLATE (unsigned int, nrow)() const noexcept

    BDENSE_TEMPLATE (unsigned int, ncol)() const noexcept

    BDENSE_TEMPLATE (unsigned int, nnozero)() const noexcept

• BDENSE TEMPLATE (Cell< Cell Type >, default val)() const
• BDENSE TEMPLATE (BDENSE TYPE() &, operator+=)(const std
• BDENSE TEMPLATE (BDENSE TYPE() &, operator-=)(const std

    BDENSE TEMPLATE (void, insert cell)(uint i

if (el[POS(i, j)]==BARRY_ZERO_DENSE)

    BDENSE_TEMPLATE (void, swap_cells)(uint i0

• if ((i0==i1) &&(j0==j1)) return
• rm cell (i0, j0, false, false)
• rm_cell (i1, j1, false, false)
• insert cell (i0, j0, val1, false, false)

    insert cell (i1, j1, val0, false, false)

• BDENSE_TEMPLATE (void, toggle_cell)(uint i
• else rm cell (i, j, false, false)
• BDENSE TEMPLATE (void, swap rows)(uint i0

    BDENSE TEMPLATE (void, swap cols)(uint j0

• BDENSE_TEMPLATE (void, zero_row)(uint i
• if (el rowsums[i]==ZERO CELL) return
• BDENSE_TEMPLATE (void, zero_col)(uint j

    if (el colsums[i]==ZERO CELL) return

    BDENSE TEMPLATE (void, transpose)()

• BDENSE TEMPLATE (void, clear)(bool hard)

    BDENSE TEMPLATE (void, resize)(uint N

• el resize (N_ *M_, ZERO_CELL)
• el rowsums resize (N, ZERO CELL)
• el colsums resize (M, ZERO CELL)
• BDENSE TEMPLATE (void, reserve)()
```

• BDENSE_TEMPLATE (void, print)(const char *fmt

va_start (args, fmt)

```
vprintf (fmt, args)
va_end (args)
BDENSE_TEMPLATE (const std::vector< Cell_Type > &, get_data)() const
BDENSE_TEMPLATE (const Cell_Type, rowsum)(unsigned int i) const
BDENSE_TEMPLATE (const Cell_Type, colsum)(unsigned int j) const
```

Variables

```
    uint M

• uint const std::vector< uint > & source
• uint const std::vector< uint > const std::vector< uint > & target

    uint const std::vector< uint > const std::vector< uint > const std::vector< Cell_Type > & value

    uint const std::vector< uint > const std::vector< cell_Type > bool add

    if(source.size() !=value.size()) throw std N = N_

    M = M_

    return

· bool copy_data
· bool delete_data_
• data = data_
delete_data = delete_data_
· uint j const
uint j
• return el [POS(i, j)] == ZERO_CELL
· return ans

    uint const Cell
    Cell_Type > & v

    uint const Cell
    Cell Type > bool check bounds

    uint const Cell
    Cell_Type > bool bool check_exists

el_rowsums [i] = (v.value - old)
• el_colsums [j] = (v.value - old)
uint j0
· uint uint i1
• uint uint uint j1
• uint uint uint bool int int * report
• Cell Type val0 = el[POS(i0,j0)]
• Cell_Type val1 = el[POS(i1,j1)]

    false

col
```

8.9.1 Macro Definition Documentation

8.9.1.1 BDENSE_TEMPLATE

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

8.9.1.2 BDENSE_TEMPLATE_ARGS

```
#define BDENSE_TEMPLATE_ARGS( ) <typename Cell_Type, typename Data_Type>
```

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

8.9.1.3 BDENSE_TYPE

```
#define BDENSE_TYPE( ) BArrayDense<Cell_Type, Data_Type>
```

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

8.9.1.4 COL

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

8.9.1.5 POS

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

Definition at line 32 of file barraydense-meat.hpp.

8.9.1.6 POS_N

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

8.9.1.7 ROW

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

8.9.1.8 ZERO_CELL

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

Definition at line 38 of file barraydense-meat.hpp.

8.9.2 Function Documentation

8.9.2.1 ans()

8.9.2.2 BDENSE_TEMPLATE() [1/37]

Definition at line 240 of file barraydense-meat.hpp.

8.9.2.3 BDENSE_TEMPLATE() [2/37]

8.9.2.4 BDENSE_TEMPLATE() [3/37]

```
BDENSE_TEMPLATE (
          BArrayDense )
```

8.9.2.5 BDENSE_TEMPLATE() [4/37]

```
BDENSE_TEMPLATE ( \sim BArrayDense )
```

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

8.9.2.6 BDENSE_TEMPLATE() [5/37]

Definition at line 558 of file barraydense-meat.hpp.

8.9.2.7 BDENSE_TEMPLATE() [6/37]

Definition at line 576 of file barraydense-meat.hpp.

8.9.2.8 BDENSE_TEMPLATE() [7/37]

Definition at line 257 of file barraydense-meat.hpp.

8.9.2.9 BDENSE_TEMPLATE() [8/37]

Definition at line 194 of file barraydense-meat.hpp.

8.9.2.10 BDENSE_TEMPLATE() [9/37]

```
BDENSE_TEMPLATE (
         bool ,
         is_empty )
```

8.9.2.11 BDENSE_TEMPLATE() [10/37]

```
BDENSE_TEMPLATE (
          bool ,
          operator = = ) const &
```

Definition at line 300 of file barraydense-meat.hpp.

8.9.2.12 BDENSE_TEMPLATE() [11/37]

Definition at line 554 of file barraydense-meat.hpp.

8.9.2.13 BDENSE_TEMPLATE() [12/37]

8.9.2.14 BDENSE_TEMPLATE() [13/37]

Definition at line 991 of file barraydense-meat.hpp.

8.9.2.15 BDENSE_TEMPLATE() [14/37]

Definition at line 986 of file barraydense-meat.hpp.

8.9.2.16 BDENSE_TEMPLATE() [15/37]

Definition at line 345 of file barraydense-meat.hpp.

8.9.2.17 BDENSE_TEMPLATE() [16/37]

Definition at line 981 of file barraydense-meat.hpp.

8.9.2.18 BDENSE_TEMPLATE() [17/37]

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

8.9.2.19 BDENSE_TEMPLATE() [18/37]

Definition at line 494 of file barraydense-meat.hpp.

8.9.2.20 BDENSE_TEMPLATE() [19/37]

```
BDENSE_TEMPLATE (
          std::vector< Cell_Type > ,
          get_row_vec )
```

8.9.2.21 BDENSE_TEMPLATE() [20/37]

```
BDENSE_TEMPLATE (
         unsigned int ,
         ncol ) const [noexcept]
```

Definition at line 540 of file barraydense-meat.hpp.

8.9.2.22 BDENSE_TEMPLATE() [21/37]

```
BDENSE_TEMPLATE (
          unsigned int ,
          nnozero ) const [noexcept]
```

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

8.9.2.23 BDENSE_TEMPLATE() [22/37]

```
BDENSE_TEMPLATE (
         unsigned int ,
         nrow ) const [noexcept]
```

Definition at line 536 of file barraydense-meat.hpp.

8.9.2.24 BDENSE_TEMPLATE() [23/37]

```
BDENSE_TEMPLATE (
     void ,
     clear )
```

Definition at line 888 of file barraydense-meat.hpp.

8.9.2.25 BDENSE_TEMPLATE() [24/37]

Definition at line 394 of file barraydense-meat.hpp.

8.9.2.26 BDENSE_TEMPLATE() [25/37]

8.9.2.27 BDENSE_TEMPLATE() [26/37]

8.9.2.28 BDENSE_TEMPLATE() [27/37]

8.9.2.29 BDENSE_TEMPLATE() [28/37]

```
BDENSE_TEMPLATE (
     void ,
     reserve )
```

Definition at line 938 of file barraydense-meat.hpp.

8.9.2.30 BDENSE_TEMPLATE() [29/37]

```
BDENSE_TEMPLATE (
           void ,
           resize )
```

8.9.2.31 BDENSE_TEMPLATE() [30/37]

8.9.2.32 BDENSE_TEMPLATE() [31/37]

8.9.2.33 BDENSE_TEMPLATE() [32/37]

8.9.2.34 BDENSE_TEMPLATE() [33/37]

```
BDENSE_TEMPLATE (
     void ,
     swap_rows )
```

8.9.2.35 BDENSE_TEMPLATE() [34/37]

8.9.2.36 BDENSE_TEMPLATE() [35/37]

```
BDENSE_TEMPLATE (
     void ,
     transpose )
```

Definition at line 860 of file barraydense-meat.hpp.

8.9.2.37 BDENSE_TEMPLATE() [36/37]

```
BDENSE_TEMPLATE (
            void ,
            zero_col )
```

8.9.2.38 BDENSE_TEMPLATE() [37/37]

```
BDENSE_TEMPLATE (
     void ,
     zero_row )
```

8.9.2.39 for()

```
for ( )
```

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

```
8.9.2.40 if() [1/4]
```

```
if (  ( {\tt i0 == i1}) \ \&\& \, ( {\tt j0 == j1}) \ )
```

8.9.2.41 if() [2/4]

Definition at line 655 of file barraydense-meat.hpp.

```
8.9.2.42 if() [3/4]
```

```
if (
    el_colsums [j] = =ZERO_CELL )
```

8.9.2.43 if() [4/4]

8.9.2.44 insert_cell() [1/2]

8.9.2.45 insert_cell() [2/2]

8.9.2.46 M()

```
bool M ( \label{eq:Array_.} \text{Array}\_. \quad \textit{M} \ )
```

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

8.9.2.47 resize() [1/6]

8.9.2.48 resize() [2/6]

8.9.2.49 resize() [3/6]

```
el resize ( \label{eq:N*M, ZERO_CELL} \mbox{N * $M$,}
```

8.9.2.50 resize() [4/6]

```
el_rowsums resize (
          N ,
          ZERO_CELL )
```

8.9.2.51 resize() [5/6]

```
el resize ( \label{eq:n_*_mass} \mbox{N\_} * \mbox{\it M\_}, \\ \mbox{ZERO\_CELL} \mbox{\ })
```

8.9.2.52 resize() [6/6]

8.9.2.53 rm_cell() [1/3]

8.9.2.54 rm_cell() [2/3]

8.9.2.55 rm_cell() [3/3]

8.9.2.56 va_end()

```
va_end (
          args )
```

8.9.2.57 va_start()

```
va_start (
          args ,
          fmt )
```

8.9.2.58 vprintf()

8.9.3 Variable Documentation

8.9.3.1 add

```
uint const std::vector< uint > const std::vector< uint > bool add

Initial value:
{
    if (source.size() != target.size())
        throw std::length_error("-source- and -target- don't match on length.")
```

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

8.9.3.2 ans

```
return ans
```

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

8.9.3.3 check_bounds

```
bool check_bounds

Initial value:
{
    if (check_bounds)
        out_of_range(i0,0u);
        out_of_range(i1,0u);
    }

for (uint j = 0u; j < M; ++j)
        std::swap(el[POS(i0, j)], el[POS(i1, j)])</pre>
```

Definition at line 646 of file barraydense-meat.hpp.

8.9.3.4 check exists

```
uint bool int check_exists
Initial value:
{
    if (check_bounds)
        out_of_range(i,j)
```

Definition at line 647 of file barraydense-meat.hpp.

8.9.3.5 col

col

Definition at line 835 of file barraydense-meat.hpp.

8.9.3.6 const

const

Initial value:

```
if (i >= N)
    throw std::range_error("The row is out of range.")
```

Definition at line 352 of file barraydense-meat.hpp.

8.9.3.7 copy_data

bool copy_data

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

8.9.3.8 data

```
data = data_
```

Definition at line 334 of file barraydense-meat.hpp.

8.9.3.9 delete_data

```
delete_data = delete_data_
```

Definition at line 335 of file barraydense-meat.hpp.

8.9.3.10 delete_data_

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

8.9.3.11 el

```
return el == ZERO_CELL
```

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

8.9.3.12 el_colsums

```
el\_colsums[j] = (v.value - old)
```

Definition at line 667 of file barraydense-meat.hpp.

8.9.3.13 el_rowsums

```
el_rowsums[i] = (v.value - old)
```

Definition at line 666 of file barraydense-meat.hpp.

8.9.3.14 else

Definition at line 662 of file barraydense-meat.hpp.

8.9.3.15 false

false

Definition at line 759 of file barraydense-meat.hpp.

8.9.3.16 i1

uint i1

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

8.9.3.17 j

j

Definition at line 365 of file barraydense-meat.hpp.

8.9.3.18 j0

uint j0

Definition at line 712 of file barraydense-meat.hpp.

8.9.3.19 j1

uint j1

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

8.9.3.20 M

 $M = M_{\underline{}}$

Definition at line 57 of file barraydense-meat.hpp.

8.9.3.21 M_

```
uint M_
Initial value:
{
    std::vector< Cell_Type > el_tmp(el)
```

Definition at line 43 of file barraydense-meat.hpp.

8.9.3.22 N

```
N = N_
```

Definition at line 56 of file barraydense-meat.hpp.

8.9.3.23 report

```
uint uint uint bool int int* report

Initial value:
{
    if (check_bounds) {
        out_of_range(i0, j0);
        out_of_range(i1, j1);
    }

    if (report != nullptr)
        (*report) = EXISTS::BOTH
```

Definition at line 716 of file barraydense-meat.hpp.

8.9.3.24 return

return

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

8.9.3.25 source

```
uint const std::vector< uint >& source
```

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

8.9.3.26 target

```
uint const std::vector< uint > const std::vector< uint >& target
```

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

8.9.3.27 v

```
uint Cell_Type v
```

Definition at line 645 of file barraydense-meat.hpp.

8.9.3.28 val0

```
Cell_Type val0 = el[POS(i0,j0)]
```

Definition at line 734 of file barraydense-meat.hpp.

8.9.3.29 val1

```
Cell_Type val1 = el[POS(i1, j1)]
```

Definition at line 735 of file barraydense-meat.hpp.

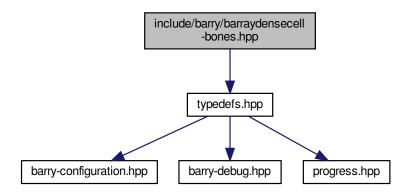
8.9.3.30 value

```
uint const std::vector< uint > const std::vector< cell_Type >&
value
```

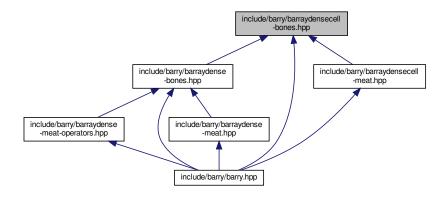
Definition at line 46 of file barraydense-meat.hpp.

8.10 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

Macros

• #define POS(a, b) (a) + (b) * N

8.10.1 Macro Definition Documentation

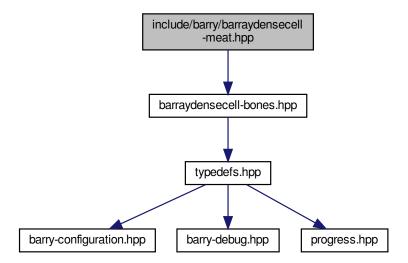
8.10.1.1 POS

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

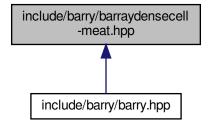
Definition at line 6 of file barraydensecell-bones.hpp.

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

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



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



Macros

```
• #define POS(a, b) (a) + (b) * dat->N
```

8.11.1 Macro Definition Documentation

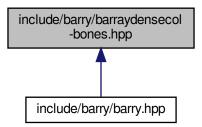
8.11.1.1 POS

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

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

8.12 include/barry/barraydensecol-bones.hpp File Reference

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



Classes

- class BArrayDenseCol< Cell_Type, Data_Type >
- class BArrayDenseCol_const< Cell_Type, Data_Type >

Macros

- #define POS(a, b) (b)*N + (a)
- #define POS_N(a, b, c) (b)*(c) + (a)
- #define ZERO_CELL static_cast<Cell_Type>(0.0)

8.12.1 Macro Definition Documentation

8.12.1.1 POS

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

Definition at line 4 of file barraydensecol-bones.hpp.

8.12.1.2 POS_N

Definition at line 5 of file barraydensecol-bones.hpp.

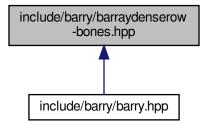
8.12.1.3 ZERO_CELL

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

Definition at line 6 of file barraydensecol-bones.hpp.

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

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



Classes

- class BArrayDenseRow< Cell_Type, Data_Type >
- class BArrayDenseRow_const< Cell_Type, Data_Type >

Macros

```
#define POS(a, b) (b) * N + (a)
#define POS_N(a, b, c) (b)*(c) + (a)
#define ZERO_CELL static_cast< Cell_Type >(0.0)
```

8.13.1 Macro Definition Documentation

8.13.1.1 POS

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

Definition at line 4 of file barraydenserow-bones.hpp.

8.13.1.2 POS N

Definition at line 5 of file barraydenserow-bones.hpp.

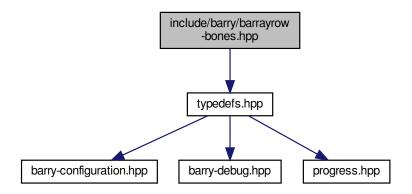
8.13.1.3 ZERO_CELL

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

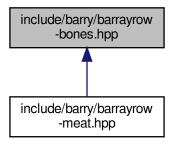
Definition at line 6 of file barraydenserow-bones.hpp.

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

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



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



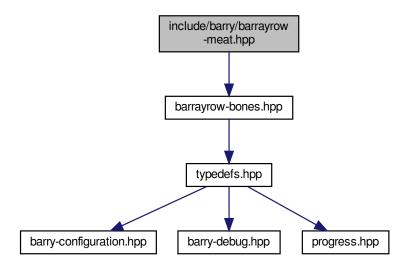
Classes

- class BArrayRow< Cell_Type, Data_Type >
- class BArrayRow_const< Cell_Type, Data_Type >

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

#include "barrayrow-bones.hpp"

Include dependency graph for barrayrow-meat.hpp:



Macros

- #define BARRY_BARRAYROW_MEAT_HPP 1
- #define BROW_TYPE() BArrayRow<Cell_Type, Data_Type>
- #define BROW_TEMPLATE_ARGS() < typename Cell_Type, typename Data_Type>
- #define BROW_TEMPLATE(a, b) template BROW_TEMPLATE_ARGS() inline a BROW_TYPE()::b

Functions

- BROW_TEMPLATE (void, operator=)(const BROW_TYPE() &val)
- BROW_TEMPLATE (void, operator+=)(const BROW_TYPE() &val)
- BROW_TEMPLATE (void, operator-=)(const BROW_TYPE() &val)
- BROW_TEMPLATE (void, operator*=)(const BROW_TYPE() &val)
- BROW TEMPLATE (void, operator/=)(const BROW TYPE() &val)

8.15.1 Macro Definition Documentation

8.15.1.1 BARRY BARRAYROW MEAT HPP

#define BARRY_BARRAYROW_MEAT_HPP 1

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

8.15.1.2 BROW_TEMPLATE

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

8.15.1.3 BROW_TEMPLATE_ARGS

```
#define BROW_TEMPLATE_ARGS() <typename Cell_Type, typename Data_Type>
```

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

8.15.1.4 BROW_TYPE

```
#define BROW_TYPE( ) BArrayRow<Cell_Type, Data_Type>
```

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

8.15.2 Function Documentation

8.15.2.1 BROW_TEMPLATE() [1/5]

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

8.15.2.2 BROW_TEMPLATE() [2/5]

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

8.15.2.3 BROW_TEMPLATE() [3/5]

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

8.15.2.4 BROW_TEMPLATE() [4/5]

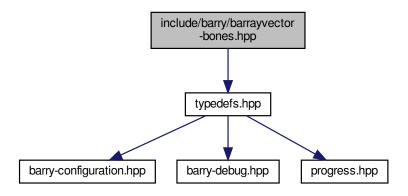
Definition at line 57 of file barrayrow-meat.hpp.

8.15.2.5 BROW_TEMPLATE() [5/5]

Definition at line 13 of file barrayrow-meat.hpp.

8.16 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:

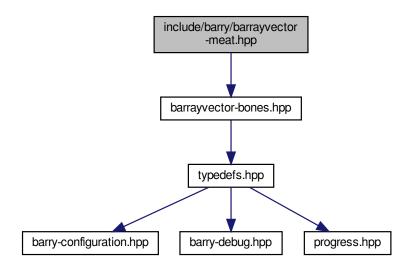


Classes

- class BArrayVector< Cell_Type, Data_Type >
 Row or column of a BArray
- class BArrayVector_const< Cell_Type, Data_Type >

8.17 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

8.17.1 Macro Definition Documentation

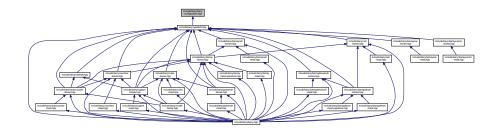
8.17.1.1 BARRY_BARRAYVECTOR_MEAT_HPP

#define BARRY_BARRAYVECTOR_MEAT_HPP 1

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

8.18 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.
- BARRY_DEBUG_LEVEL, when defined, will make things verbose.
- #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< size_t >(UINT_MAX/2u)
- template<typename Ta , typename Tb >
 using Map = std::map< Ta, Tb >

8.18.1 Macro Definition Documentation

8.18.1.1 BARRY_CHECK_SUPPORT

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

Definition at line 47 of file barry-configuration.hpp.

8.18.1.2 BARRY_ISFINITE

Definition at line 40 of file barry-configuration.hpp.

8.18.1.3 BARRY_MAX_NUM_ELEMENTS

```
#define BARRY_MAX_NUM_ELEMENTS static_cast< size_t > (UINT_MAX/2u)
```

Definition at line 55 of file barry-configuration.hpp.

8.18.1.4 BARRY_SAFE_EXP

```
#define BARRY_SAFE_EXP -100.0
```

Definition at line 33 of file barry-configuration.hpp.

8.18.1.5 printf_barry

```
#define printf_barry printf
```

Definition at line 51 of file barry-configuration.hpp.

8.18.2 Typedef Documentation

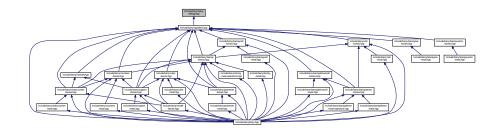
8.18.2.1 Map

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

Definition at line 27 of file barry-configuration.hpp.

8.19 include/barry/barry-debug.hpp File Reference

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



Macros

#define BARRY_DEBUG_LEVEL 0

8.19.1 Macro Definition Documentation

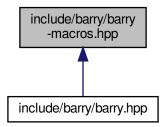
8.19.1.1 BARRY_DEBUG_LEVEL

#define BARRY_DEBUG_LEVEL 0

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

8.20 include/barry/barry-macros.hpp File Reference

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



Macros

- #define BARRY_ZERO Cell<Cell_Type>(0.0)
- #define BARRY_ZERO_DENSE static_cast<Cell_Type>(0.0)
- #define BARRY_ONE Cell<Cell_Type>(1.0)
- #define BARRY_ONE_DENSE static_cast<Cell_Type>(1.0)
- #define BARRY_UNUSED(expr) do { (void)(expr); } while (0);

8.20.1 Macro Definition Documentation

8.20.1.1 BARRY_ONE

```
#define BARRY_ONE CellCell_Type>(1.0)
```

Definition at line 7 of file barry-macros.hpp.

8.20.1.2 BARRY_ONE_DENSE

```
#define BARRY_ONE_DENSE static_cast<Cell_Type>(1.0)
```

Definition at line 8 of file barry-macros.hpp.

8.20.1.3 BARRY_UNUSED

Definition at line 10 of file barry-macros.hpp.

8.20.1.4 BARRY_ZERO

```
#define BARRY_ZERO Cell<Cell_Type>(0.0)
```

Definition at line 4 of file barry-macros.hpp.

8.20.1.5 BARRY_ZERO_DENSE

```
#define BARRY_ZERO_DENSE static_cast<Cell_Type>(0.0)
```

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

8.21 include/barry/barry.hpp File Reference

```
#include <iostream>
#include <cstdarg>
#include <vector>
#include <unordered_map>
#include <functional>
#include <stdexcept>
#include <cmath>
#include <map>
#include <algorithm>
#include <utility>
#include <random>
#include <climits>
#include <cfloat>
#include <string>
#include <cstdint>
#include "typedefs.hpp"
#include "barry-macros.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 "barraydense-bones.hpp"
#include "barraydensecell-bones.hpp"
#include "barraydenserow-bones.hpp"
#include "barraydensecol-bones.hpp"
#include "barraydense-meat.hpp"
#include "barraydensecell-meat.hpp"
#include "barraydense-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)

8.21.1 Macro Definition Documentation

8.21.1.1 BARRY_HPP

#define BARRY_HPP

Definition at line 22 of file barry.hpp.

8.21.1.2 BARRY_VERSION

#define BARRY_VERSION 0.1

Definition at line 24 of file barry.hpp.

8.21.1.3 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 86 of file barry.hpp.

8.21.1.4 COUNTER_LAMBDA

Value:

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

Definition at line 89 of file barry.hpp.

8.21.1.5 RULE_FUNCTION

Value:

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

Definition at line 93 of file barry.hpp.

8.21.1.6 **RULE LAMBDA**

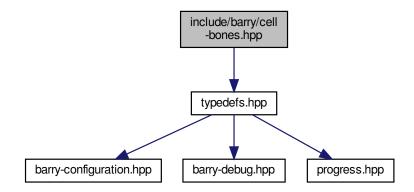
Value:

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

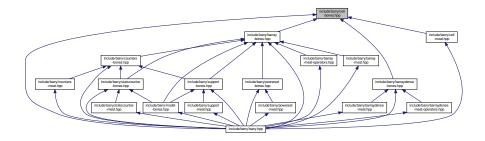
Definition at line 96 of file barry.hpp.

8.22 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:



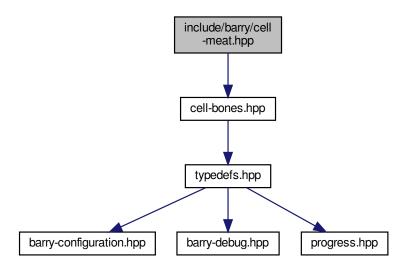
Classes

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

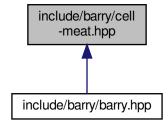
8.23 include/barry/cell-meat.hpp File Reference

#include "cell-bones.hpp"

Include dependency graph for cell-meat.hpp:



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



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

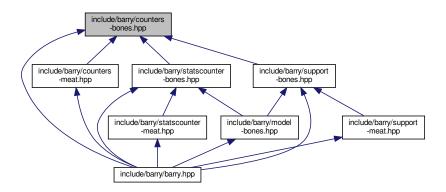
8.25 include/barry/counters-bones.hpp File Reference

```
#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.

8.26 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:



Macros

- #define COUNTER_TYPE() Counter<Array_Type,Data_Type>
- #define COUNTER_TEMPLATE_ARGS() < typename Array_Type, typename Data_Type >

- #define COUNTERS_TYPE() Counters<Array_Type,Data_Type>
- #define COUNTERS_TEMPLATE_ARGS() < typename Array_Type, typename Data_Type >

Functions

- COUNTER_TEMPLATE (, Counter)(const Counter< Array_Type
- Data Type init fun (counter .init fun)
- Data Type &&counter init fun (std::move(counter .init fun))
- Data_Type &&counter_ data (std::move(counter_.data))
- Data_Type &&counter_ name (std::move(counter_.name))
- Data_Type &&counter_ desc (std::move(counter_.desc))

Move constructor.

- COUNTER TEMPLATE (COUNTER TYPE(), operator=)(const Counter< Array Type
- COUNTER_TEMPLATE (COUNTER_TYPE() &, operator=)(Counter< Array_Type
- COUNTER TEMPLATE (double, count)(Array Type & Array

< Move assignment

- return count_fun (Array, i, j, data)
- COUNTER TEMPLATE (double, init)(Array Type & Array
- return init_fun (Array, i, j, data)
- COUNTER_TEMPLATE (std::string, get_name)() const
- COUNTER_TEMPLATE (std::string, get_description)() const
- COUNTERS_TEMPLATE (, Counters)()
- COUNTERS_TEMPLATE (COUNTER_TYPE() &, operator[])(uint idx)
- Data_Type Counters (Counters < Array_Type, Data_Type > &&counters_) noexcept
- COUNTERS_TEMPLATE (COUNTERS_TYPE(), operator=)(const Counters < Array_Type
- COUNTERS_TEMPLATE (COUNTERS_TYPE() &, operator=)(Counters< Array_Type
- COUNTERS TEMPLATE (void, add counter)(Counter< Array Type
- COUNTERS_TEMPLATE (std::vector< std::string >, get_names)() const
- COUNTERS_TEMPLATE (std::vector< std::string >, get_descriptions)() const

Variables

- Data Type & counter
- Data_Type &&counter_ noexcept
- uint i
- uint uint j
- return * this
- Data_Type counter
- return
- Data_Type count_fun_
- Data_Type Counter_fun_type
 Array_Type, Data_Type > init_fun_
- Data_Type Counter_fun_type
 Array_Type, Data_Type > Data_Type data_
- Data_Type Counter_fun_type
 Array_Type, Data_Type > Data_Type std::string name_
- Data_Type Counter_fun_type
 Array_Type, Data_Type > Data_Type std::string std::string desc_

8.26.1 Macro Definition Documentation

8.26.1.1 COUNTER_TEMPLATE

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

8.26.1.2 COUNTER_TEMPLATE_ARGS

```
#define COUNTER_TEMPLATE_ARGS() <typename Array_Type, typename Data_Type>
```

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

8.26.1.3 COUNTER_TYPE

```
#define COUNTER_TYPE( ) Counter<Array_Type, Data_Type>
```

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

8.26.1.4 COUNTERS_TEMPLATE

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

8.26.1.5 COUNTERS_TEMPLATE_ARGS

```
#define COUNTERS_TEMPLATE_ARGS() <typename Array_Type, typename Data_Type>
```

Definition at line 116 of file counters-meat.hpp.

8.26.1.6 COUNTERS_TYPE

```
#define COUNTERS_TYPE( ) Counters<Array_Type,Data_Type>
```

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

8.26.2 Function Documentation

8.26.2.1 count fun()

8.26.2.2 COUNTER_TEMPLATE() [1/7]

```
COUNTER_TEMPLATE (

Counter ) const
```

8.26.2.3 COUNTER_TEMPLATE() [2/7]

8.26.2.4 COUNTER_TEMPLATE() [3/7]

8.26.2.5 COUNTER_TEMPLATE() [4/7]

< Move assignment

8.26.2.6 COUNTER_TEMPLATE() [5/7]

```
COUNTER_TEMPLATE ( \label{eq:counter} \mbox{double ,} \\ \mbox{init ) } \&
```

8.26.2.7 COUNTER_TEMPLATE() [6/7]

```
COUNTER_TEMPLATE (
          std::string ,
          get_description ) const
```

Definition at line 106 of file counters-meat.hpp.

8.26.2.8 COUNTER_TEMPLATE() [7/7]

```
COUNTER_TEMPLATE (
          std::string ,
          get_name ) const
```

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

8.26.2.9 Counters()

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

8.26.2.10 COUNTERS_TEMPLATE() [1/7]

```
COUNTERS_TEMPLATE (

Counters )
```

Definition at line 121 of file counters-meat.hpp.

8.26.2.11 COUNTERS_TEMPLATE() [2/7]

Definition at line 123 of file counters-meat.hpp.

8.26.2.12 COUNTERS_TEMPLATE() [3/7]

```
COUNTERS_TEMPLATE (

COUNTERS_TYPE() & ,

operator )
```

8.26.2.13 COUNTERS_TEMPLATE() [4/7]

```
COUNTERS_TEMPLATE (

COUNTERS_TYPE() ,

operator ) const
```

8.26.2.14 COUNTERS_TEMPLATE() [5/7]

```
COUNTERS_TEMPLATE (
          std::vector< std::string > ,
          get_descriptions ) const
```

Definition at line 194 of file counters-meat.hpp.

8.26.2.15 COUNTERS_TEMPLATE() [6/7]

```
COUNTERS_TEMPLATE (
          std::vector< std::string > ,
          get_names ) const
```

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

8.26.2.16 COUNTERS_TEMPLATE() [7/7]

```
COUNTERS_TEMPLATE (
     void ,
     add_counter )
```

8.26.2.17 data()

8.26.2.18 desc()

Move constructor.

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

8.26.2.19 init_fun() [1/3]

8.26.2.20 init_fun() [2/3]

Definition at line 15 of file counters-meat.hpp.

8.26.2.21 init_fun() [3/3]

8.26.2.22 name()

8.26.3 Variable Documentation

8.26.3.1 count_fun_

```
Data_Type count_fun_
```

Definition at line 163 of file counters-meat.hpp.

8.26.3.2 counter

```
Data_Type counter

Initial value:
{
    data.push_back(counter)
```

Definition at line 154 of file counters-meat.hpp.

8.26.3.3 counter_

return *this

```
Data_Type & counter_

Initial value:
{
    if (this != &counter_) {
        this->count_fun = counter_.count_fun;
        this->init_fun = counter_.init_fun;

        this->data = counter_.data;
        this->name = counter_.name;
        this->desc = counter_.desc;
}
```

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

8.26.3.4 data_

```
Data_Type Counter_fun_type<Array_Type,Data_Type> Data_Type data_
```

Definition at line 165 of file counters-meat.hpp.

8.26.3.5 desc

```
Data_Type Counter_fun_type<Array_Type,Data_Type > Data_Type std::string std::string desc_
```

Initial value:

```
[
```

```
data.push_back(Counter<Array_Type,Data_Type>(
    count_fun_,
    init_fun_,
    data_,
    name_,
    desc_
```

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

8.26.3.6 i

uint i

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

8.26.3.7 init_fun_

```
Data_Type Counter_fun_type<Array_Type,Data_Type> init_fun_
```

Definition at line 164 of file counters-meat.hpp.

8.26.3.8 j

```
uint uint j
```

Initial value:

```
if (count_fun == nullptr)
    return 0.0
```

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

8.26.3.9 name_

```
Data_Type Counter_fun_type<Array_Type,Data_Type> Data_Type std::string name_
```

Definition at line 166 of file counters-meat.hpp.

8.26.3.10 noexcept

```
Data_Type &&counters_ noexcept
```

Initial value:

```
if (this != &counter_)
{
    this->data = std::move(counter_.data);

    this->count_fun = std::move(counter_.count_fun);
    this->init_fun = std::move(counter_.init_fun);

    this->name = std::move(counter_.name);
    this->desc = std::move(counter_.desc);
}
return *this
```

Definition at line 28 of file counters-meat.hpp.

8.26.3.11 return

return

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

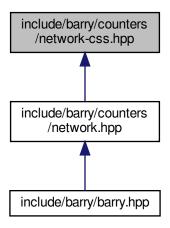
8.26.3.12 this

```
return* this
```

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

8.27 include/barry/counters/network-css.hpp File Reference

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



Macros

- #define CSS_SIZE()
- #define CSS_CASE_TRUTH() if ((i < n) && (j < n))
- #define CSS_TRUE_CELLS()
- #define CSS_CASE_PERCEIVED() else if ((($i \ge s$) && (i < e)) & (($j \ge s$) && (j < e)))
- #define CSS PERCEIVED CELLS()
- #define CSS CASE ELSE()
- #define CSS_CHECK_SIZE_INIT()
- #define CSS_CHECK_SIZE()
- #define CSS_APPEND(name)
- #define CSS_NET_COUNTER_LAMBDA_INIT()

Functions

• template<typename Tnet = Network> void counter_css_partially_false_recip_commi (NetCounters< Tnet > *counters, uint netsize, const std ← ::vector< uint > &end_)

Counts errors of commission.

template<typename Tnet = Network>
 void counter_css_partially_false_recip_omiss (NetCounters< Tnet > *counters, uint netsize, const std
 ::vector< uint > &end_)

Counts errors of omission.

• template<typename Tnet = Network> void counter_css_completely_false_recip_comiss (NetCounters< Tnet > *counters, uint netsize, const std ← ::vector< uint > &end_)

Counts completely false reciprocity (comission)

```
• template<typename Tnet = Network>
  void counter_css_completely_false_recip_omiss (NetCounters< Tnet > *counters, uint netsize, const std↔
  ::vector< uint > &end_)
     Counts completely false reciprocity (omission)
• template<typename Tnet = Network>
  void counter css mixed recip (NetCounters< Tnet > *counters, uint netsize, const std::vector< uint >
  &end )
     Counts mixed reciprocity errors.
• template<typename Tnet = Network>
  void counter_css_census01 (NetCounters< Tnet > *counters, uint netsize, const std::vector< uint > &end↔
• template<typename Tnet = Network>
 void counter css census02 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
template<typename Tnet = Network>
 void counter_css_census03 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
template<typename Tnet = Network>
  void counter_css_census04 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
 _)
• template<typename Tnet = Network>
 void counter css census05 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
template<typename Tnet = Network>
  void counter css census06 (NetCounters< Tnet > *counters, uint netsize, const std::vector< uint > &end↔
• template<typename Tnet = Network>
 void counter_css_census07 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
 _)
• template<typename Tnet = Network>
  void counter_css_census08 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
 _)
• template<typename Tnet = Network>
 void counter css census09 (NetCounters < Tnet > *counters, uint netsize, const std::vector < uint > &end ←
template<typename Tnet = Network>
  void counter_css_census10 (NetCounters< Tnet > *counters, uint netsize, const std::vector< uint > &end↔
  _)
```

8.27.1 Macro Definition Documentation

8.27.1.1 CSS_APPEND

Definition at line 42 of file network-css.hpp.

8.27.1.2 CSS_CASE_ELSE

```
#define CSS_CASE_ELSE( )
```

Definition at line 27 of file network-css.hpp.

8.27.1.3 CSS_CASE_PERCEIVED

```
      \# define \ CSS\_CASE\_PERCEIVED ( ) \ else \ if \ (((i >= s) \ \&\& \ (i < e)) \ \& \ ((j >= s) \ \&\& \ (j < e)))
```

Definition at line 20 of file network-css.hpp.

8.27.1.4 CSS CASE TRUTH

```
#define CSS_CASE_TRUTH( ) if ((i < n) && (j < n))
```

Definition at line 13 of file network-css.hpp.

8.27.1.5 CSS_CHECK_SIZE

Definition at line 37 of file network-css.hpp.

8.27.1.6 CSS CHECK SIZE INIT

throw std::range_error("The network does not match the prescribed size.");

Definition at line 31 of file network-css.hpp.

8.27.1.7 CSS_NET_COUNTER_LAMBDA_INIT

Definition at line 49 of file network-css.hpp.

8.27.1.8 CSS_PERCEIVED_CELLS

```
#define CSS_PERCEIVED_CELLS()

Value:
    double tji = static_cast<double>(Array(j - s, i - s, false)); \
    double pji = static_cast<double>(Array(j, i, false)); \
    double tij = static_cast<double>(Array(i - s, j - s, false));
```

Definition at line 21 of file network-css.hpp.

8.27.1.9 CSS_SIZE

```
#define CSS_SIZE()

Value:
    uint n = data.indices[0u]; \
    uint s = data.indices[1u]; \
    uint e = data.indices[2u];
```

Definition at line 7 of file network-css.hpp.

8.27.1.10 CSS_TRUE_CELLS

```
#define CSS_TRUE_CELLS()
Value:

double tji = static_cast<double>(Array(j, i, false)); \
double pij = static_cast<double>(Array(i + s, j + s, false)); \
double pji = static_cast<double>(Array(j + s, i + s, false));
```

Definition at line 14 of file network-css.hpp.

8.27.2 Function Documentation

8.27.2.1 counter_css_census01()

Definition at line 275 of file network-css.hpp.

8.27.2.2 counter_css_census02()

Definition at line 325 of file network-css.hpp.

8.27.2.3 counter_css_census03()

Definition at line 364 of file network-css.hpp.

8.27.2.4 counter_css_census04()

Definition at line 403 of file network-css.hpp.

8.27.2.5 counter_css_census05()

Definition at line 442 of file network-css.hpp.

8.27.2.6 counter_css_census06()

Definition at line 481 of file network-css.hpp.

8.27.2.7 counter_css_census07()

Definition at line 520 of file network-css.hpp.

8.27.2.8 counter_css_census08()

Definition at line 559 of file network-css.hpp.

8.27.2.9 counter_css_census09()

Definition at line 598 of file network-css.hpp.

8.27.2.10 counter_css_census10()

Definition at line 637 of file network-css.hpp.

8.27.2.11 counter_css_completely_false_recip_comiss()

Counts completely false reciprocity (comission)

Definition at line 154 of file network-css.hpp.

8.27.2.12 counter_css_completely_false_recip_omiss()

Counts completely false reciprocity (omission)

Definition at line 194 of file network-css.hpp.

8.27.2.13 counter_css_mixed_recip()

Counts mixed reciprocity errors.

Definition at line 234 of file network-css.hpp.

8.27.2.14 counter_css_partially_false_recip_commi()

Counts errors of commission.

Parameters

netsize	Size of the reference (true) network
end←	Vector indicating one past the ending index of each network. (see details)
_	

The end_ parameter should be of length ${\tt N}$ of networks - 1. It is assumed that the first network ends at netsize.

Definition at line 63 of file network-css.hpp.

8.27.2.15 counter_css_partially_false_recip_omiss()

Counts errors of omission.

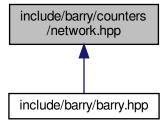
Definition at line 110 of file network-css.hpp.

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

#include "network-css.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)
- #define NETWORKDENSE_COUNTER_LAMBDA(a)

Macros for defining rules

- #define NETWORK RULE(a)
- #define NETWORK_RULE_LAMBDA(a)

Functions

```
template<typename Tnet = Network>
  void counter_edges (NetCounters < Tnet > *counters)
     Number of edges.
• template<typename Tnet = Network>
  void counter isolates (NetCounters < Tnet > *counters)
     Number of isolated vertices.

    template<> void counter_isolates (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter mutual (NetCounters< Tnet > *counters)
     Number of mutual ties.
• template<typename Tnet = Network>
  void counter_istar2 (NetCounters < Tnet > *counters)

    template<> void counter_istar2 (NetCounters< NetworkDense > *counters)

    template<typename Tnet = Network>

  void counter ostar2 (NetCounters< Tnet > *counters)

    template<> void counter ostar2 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter_ttriads (NetCounters < Tnet > *counters)

    template<> void counter_ttriads (NetCounters< NetworkDense > *counters)

template<typename Tnet = Network>
  void counter ctriads (NetCounters< Tnet > *counters)

    template<> void counter ctriads (NetCounters< NetworkDense > *counters)

template<typename Tnet = Network>
  void counter_density (NetCounters < Tnet > *counters)
• template<typename Tnet = Network>
  void counter_idegree15 (NetCounters < Tnet > *counters)

    template<> void counter idegree15 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter odegree15 (NetCounters< Tnet > *counters)

    template<> void counter_odegree15 (NetCounters< NetworkDense > *counters)

• template<typename Tnet = Network>
  void counter_absdiff (NetCounters< Tnet > *counters, uint attr_id, double alpha=1.0)
     Sum of absolute attribute difference between ego and alter.
• template<typename Tnet = Network>
  void counter diff (NetCounters < Tnet > *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)
```

• template<typename Tnet = Network>

• template<typename Tnet = Network>

void counter_nodeicov (NetCounters< Tnet > *counters, uint attr_id)

void counter_nodeocov (NetCounters< Tnet > *counters, uint attr_id)

```
    template<typename Tnet = Network>
        void counter_nodecov (NetCounters< Tnet > *counters, uint attr_id)
    template<typename Tnet = Network>
        void counter_nodematch (NetCounters< Tnet > *counters, uint attr_id)
    template<typename Tnet = Network>
        void counter_idegree (NetCounters< Tnet > *counters, std::vector< uint > d)
    Counts number of vertices with a given in-degree.
    template<>> void counter_idegree (NetCounters< NetworkDense > *counters, std::vector< uint > d)
    template<typename Tnet = Network>
        void counter_odegree (NetCounters< Tnet > *counters, std::vector< uint > d)
    Counts number of vertices with a given out-degree.
    template<>> void counter_odegree (NetCounters<< NetworkDense > *counters, std::vector< uint > d)
    template<typename Tnet = Network>
        void counter_degree (NetCounters< Tnet > *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 >).

template < typename Tnet = Network >
 void rules_zerodiag (NetRules < Tnet > *rules)
 Number of edges.

Convenient typedefs for network objects.

- #define BARRY_ZERO_NETWORK 0.0
- #define BARRY_ZERO_NETWORK_DENSE 0
- typedef BArray< double, NetworkData > Network
- typedef BArrayDense< int, NetworkData > NetworkDense
- template<typename Tnet = Network>
 using NetCounter = Counter< Tnet, NetCounterData >
- template<typename Tnet = Network>
 using NetCounters = Counters< Tnet, NetCounterData >
- template<typename Tnet = Network>
 using NetSupport = Support< Tnet, NetCounterData >
- template < typename Tnet = Network >
 using NetStatsCounter = StatsCounter < Tnet, NetCounterData >
- template < typename Tnet >
 using NetModel = Model < Tnet, NetCounterData >
- template<typename Tnet = Network>
 using NetRule = Rule
 Tnet, bool >
- template<typename Tnet = Network>
 using NetRules = Rules< Tnet, bool >

8.28.1 Macro Definition Documentation

8.28.1.1 BARRY_ZERO_NETWORK

```
#define BARRY_ZERO_NETWORK 0.0
```

Definition at line 85 of file network.hpp.

8.28.1.2 BARRY_ZERO_NETWORK_DENSE

```
#define BARRY_ZERO_NETWORK_DENSE 0
```

Definition at line 86 of file network.hpp.

8.28.1.3 NET_C_DATA_IDX

Definition at line 74 of file network.hpp.

8.28.1.4 NET_C_DATA_NUM

```
\label{eq:continuous} \begin{tabular}{ll} \# define & NET_C_DATA_NUM( & & & \\ & i & ) & (data.numbers[i]) \\ \end{tabular}
```

Definition at line 75 of file network.hpp.

8.28.1.5 NETWORK_COUNTER

Value:

```
template<typename Tnet = Network>\
inline double (a) (const Tnet & Array, uint i, uint j, NetCounterData & data)
```

Function for definition of a network counter function

Definition at line 114 of file network.hpp.

8.28.1.6 NETWORK_COUNTER_LAMBDA

Lambda function for definition of a network counter function

Definition at line 119 of file network.hpp.

8.28.1.7 NETWORK_RULE

Function for definition of a network counter function

Definition at line 133 of file network.hpp.

8.28.1.8 NETWORK_RULE_LAMBDA

Lambda function for definition of a network counter function

Definition at line 138 of file network.hpp.

8.28.1.9 NETWORKDENSE_COUNTER_LAMBDA

Definition at line 123 of file network.hpp.

8.28.2 Typedef Documentation

8.28.2.1 NetCounter

```
template<typename Tnet = Network>
using NetCounter = Counter<Tnet, NetCounterData >
```

Definition at line 89 of file network.hpp.

8.28.2.2 NetCounters

```
template<typename Tnet = Network>
using NetCounters = Counters<Tnet, NetCounterData>
```

Definition at line 92 of file network.hpp.

8.28.2.3 NetModel

```
template<typename Tnet >
using NetModel = Model<Tnet, NetCounterData>
```

Definition at line 101 of file network.hpp.

8.28.2.4 NetRule

```
template<typename Tnet = Network>
using NetRule = Rule<Tnet, bool>
```

Definition at line 104 of file network.hpp.

8.28.2.5 NetRules

```
template<typename Tnet = Network>
using NetRules = Rules<Tnet, bool>
```

Definition at line 107 of file network.hpp.

8.28.2.6 NetStatsCounter

```
template<typename Tnet = Network>
using NetStatsCounter = StatsCounter<Tnet, NetCounterData>
```

Definition at line 98 of file network.hpp.

8.28.2.7 NetSupport

```
template<typename Tnet = Network>
using NetSupport = Support<Tnet, NetCounterData >
```

Definition at line 95 of file network.hpp.

8.28.2.8 Network

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

Definition at line 82 of file network.hpp.

8.28.2.9 NetworkDense

```
typedef BArrayDense<int, NetworkData> NetworkDense
```

Definition at line 83 of file network.hpp.

8.28.3 Function Documentation

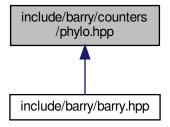
8.28.3.1 rules_zerodiag()

Number of edges.

Definition at line 1377 of file network.hpp.

8.29 include/barry/counters/phylo.hpp File Reference

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



Classes

- · class NodeData
 - Data definition for the PhyloArray class.
- · class PhyloCounterData
- class PhyloRuleDynData

Macros

- #define DEFAULT_DUPLICATION 1u
- #define DUPL_SPEC 0u
- #define DUPL_DUPL 1u
- #define DUPL_EITH 2u
- #define MAKE_DUPL_VARS()
- #define IS_EITHER() (DATA_AT == DUPL_EITH)
- #define IS_DUPLICATION() ((DATA_AT == DUPL_DUPL) & (DPL))
- #define IS_SPECIATION() ((DATA_AT == DUPL_SPEC) & (!DPL))
- #define IF_MATCHES()
- #define IF_NOTMATCHES()
- #define PHYLO_COUNTER_LAMBDA(a)

Extension of a simple counter.

- #define PHYLO_RULE_DYN_LAMBDA(a)
- #define PHYLO_CHECK_MISSING()

Typedefs

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

Convenient typedefs for Node objects.

- typedef BArrayDense< 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 (unsigned int d)
- void counter_overall_gains (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)
 Overall functional gains.
- void counter_gains (PhyloCounters *counters, std::vector< uint > nfun, unsigned int duplication=DEFAULT_DUPLICATION)

 Functional gains for a specific function (nfun).
- void counter_gains_k_offspring (PhyloCounters *counters, std::vector< uint > nfun, uint k=1u, unsigned int duplication=DEFAULT_DUPLICATION)

k genes gain function nfun

- void counter_genes_changing (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)
- void counter_preserve_pseudogene (PhyloCounters *counters, unsigned int nfunA, unsigned int nfunB, unsigned int duplication=DEFAULT DUPLICATION)

Keeps track of how many pairs of genes preserve pseudostate.

- void counter_prop_genes_changing (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Keeps track of how many genes are changing (either 0, 1, or 2 if dealing with regular trees.)
- void counter_overall_loss (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)
 Overall functional loss.
- void counter_maxfuns (PhyloCounters *counters, uint lb, uint ub, unsigned int duplication=DEFAULT_DUPLICATION)

 Cap the number of functions per gene.
- void counter_loss (PhyloCounters *counters, std::vector < uint > nfun, unsigned int duplication=DEFAULT_DUPLICATION)

 Total count of losses for an specific function.
- void counter_overall_changes (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

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

 Total count of Sub-functionalization events.
- void counter_cogain (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

 Co-evolution (joint gain or loss)
- void counter_longest (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

Longest branch mutates (either by gain or by loss)

- void counter_neofun (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

 Total number of neofunctionalization events.
- void counter_pairwise_neofun_singlefun (PhyloCounters *counters, uint nfunA, unsigned int duplication=DEFAULT_DUPLICATI

 Total number of neofunctionalization events sum_u sum_{w < u} [x(u,a)*(1 x(w,a)) + (1 x(u,a)) * x(w,a)] change

 stat: delta{x(u,a): 0->1} = 1 2 * x(w,a)
- void counter_neofun_a2b (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION

 Total number of neofunctionalization events.
- void counter_co_opt (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

Function co-opting.

- void counter_k_genes_changing (PhyloCounters *counters, unsigned int k, unsigned int duplication=DEFAULT_DUPLICATION)
 Indicator function. Equals to one if k genes changed and zero otherwise.
- void counter_less_than_p_prop_genes_changing (PhyloCounters *counters, double p, unsigned int duplication=DEFAULT DUPLICATION)

Indicator function. Equals to one if k genes changed and zero otherwise.

- void counter_gains_from_0 (PhyloCounters *counters, std::vector< uint > nfun, unsigned int duplication=DEFAULT_DUPLICAT

 Used when all the functions are in 0 (like the root node prob.)
- void counter_overall_gains_from_0 (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Used when all the functions are in 0 (like the root node prob.)
- void counter_pairwise_overall_change (PhyloCounters *counters, unsigned int duplication=DEFAULT_DUPLICATION)

 Used when all the functions are in 0 (like the root node prob.)
- void counter_pairwise_preserving (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT_DUPLICATION)

Used when all the functions are in 0 (like the root node prob.)

 void counter_pairwise_first_gain (PhyloCounters *counters, uint nfunA, uint nfunB, unsigned int duplication=DEFAULT DUPLICATION)

Used when all the functions are in 0 (like the root node prob.)

void rule_dyn_limit_changes (PhyloSupport *support, uint pos, uint lb, uint ub, unsigned int duplication=DEFAULT_DUPLICATIC
 Overall functional gains.

8.29.1 Macro Definition Documentation

8.29.1.1 DEFAULT_DUPLICATION

#define DEFAULT_DUPLICATION 1u

Definition at line 5 of file phylo.hpp.

8.29.1.2 DUPL_DUPL

#define DUPL_DUPL 1u

Definition at line 7 of file phylo.hpp.

8.29.1.3 DUPL EITH

#define DUPL_EITH 2u

Definition at line 8 of file phylo.hpp.

8.29.1.4 DUPL_SPEC

```
#define DUPL_SPEC Ou
```

Definition at line 6 of file phylo.hpp.

8.29.1.5 IF_MATCHES

```
#define IF_MATCHES()

Value:
    MAKE_DUPL_VARS() \
    if (IS_EITHER() | IS_DUPLICATION() | IS_SPECIATION())
```

Definition at line 19 of file phylo.hpp.

8.29.1.6 IF_NOTMATCHES

```
#define IF_NOTMATCHES( )

Value:
    MAKE_DUPL_VARS() \
    if (!IS_EITHER() & !IS_DUPLICATION() & !IS_SPECIATION())
```

Definition at line 21 of file phylo.hpp.

8.29.1.7 IS_DUPLICATION

```
#define IS_DUPLICATION() ((DATA_AT == DUPL_DUPL) & (DPL))
```

Definition at line 16 of file phylo.hpp.

8.29.1.8 IS_EITHER

```
#define IS_EITHER( ) (DATA_AT == DUPL_EITH)
```

Definition at line 15 of file phylo.hpp.

8.29.1.9 IS_SPECIATION

```
#define IS_SPECIATION( ) ((DATA_AT == DUPL_SPEC) & (!DPL))
```

Definition at line 17 of file phylo.hpp.

8.29.1.10 MAKE_DUPL_VARS

```
#define MAKE_DUPL_VARS( )

Value:
   bool DPL = Array.D()->duplication; \
   unsigned int DATA_AT = data[Ou];
```

Definition at line 11 of file phylo.hpp.

8.29.1.11 PHYLO_CHECK_MISSING

```
#define PHYLO_CHECK_MISSING( )

Value:
    if (Array.D() == nullptr) \
    throw std::logic_error("The array data is nullptr."); \
```

Definition at line 139 of file phylo.hpp.

8.29.1.12 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 133 of file phylo.hpp.

8.29.1.13 PHYLO_RULE_DYN_LAMBDA

Value:

```
Rule_fun_type<PhyloArray, PhyloRuleDynData> a = \
[](const PhyloArray & Array, uint i, uint j, PhyloRuleDynData & data)
```

Definition at line 136 of file phylo.hpp.

8.29.2 Typedef Documentation

8.29.2.1 PhyloArray

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

Definition at line 106 of file phylo.hpp.

8.29.2.2 PhyloCounter

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

Definition at line 107 of file phylo.hpp.

8.29.2.3 PhyloCounters

```
typedef Counters< PhyloArray, PhyloCounterData> PhyloCounters
```

Definition at line 108 of file phylo.hpp.

8.29.2.4 PhyloModel

 ${\tt typedef\ Model < PhyloArray,\ PhyloCounterData,\ PhyloRuleData,\ PhyloRuleDynData > PhyloModel}$

Definition at line 118 of file phylo.hpp.

8.29.2.5 PhyloPowerSet

typedef PowerSet<PhyloArray, PhyloRuleData> PhyloPowerSet

Definition at line 119 of file phylo.hpp.

8.29.2.6 PhyloRule

typedef Rule<PhyloArray,PhyloRuleData> PhyloRule

Definition at line 110 of file phylo.hpp.

8.29.2.7 PhyloRuleData

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

Definition at line 99 of file phylo.hpp.

8.29.2.8 PhyloRuleDyn

typedef Rule<PhyloArray,PhyloRuleDynData> PhyloRuleDyn

Definition at line 113 of file phylo.hpp.

8.29.2.9 PhyloRules

typedef Rules<PhyloArray,PhyloRuleData> PhyloRules

Definition at line 111 of file phylo.hpp.

8.29.2.10 PhyloRulesDyn

typedef Rules<PhyloArray,PhyloRuleDynData> PhyloRulesDyn

Definition at line 114 of file phylo.hpp.

8.29.2.11 PhyloStatsCounter

typedef StatsCounter<PhyloArray, PhyloCounterData> PhyloStatsCounter Definition at line 117 of file phylo.hpp.

8.29.2.12 PhyloSupport

typedef Support<PhyloArray, PhyloCounterData, PhyloRuleData, PhyloRuleDynData > PhyloSupport
Definition at line 116 of file phylo.hpp.

8.29.3 Function Documentation

8.29.3.1 get_last_name()

```
\begin{tabular}{lll} {\tt std::string get\_last\_name (} \\ & {\tt unsigned int } d \end{tabular} ) & [inline] \\ \end{tabular}
```

Definition at line 142 of file phylo.hpp.

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

```
#include "barray-bones.hpp"
#include "support-bones.hpp"
#include "statscounter-bones.hpp"
#include "rules-bones.hpp"
Include dependency graph for model-bones.hpp:
```

support-bones.hpp statscounter-bones.hpp

rules-bones.hpp counters-bones.hpp

barray-bones.hpp

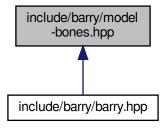
typedefs.hpp

barry-configuration.hpp

barry-debug.hpp

progress.hpp

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



Classes

class Model < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >

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

Functions

template<typename Array_Type >
 std::vector< double > keygen_default (const Array_Type &Array_)
 Array Hasher class (used for computing support)

8.30.1 Function Documentation

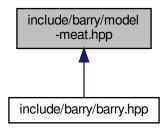
8.30.1.1 keygen_default()

Array Hasher class (used for computing support)

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

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

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



Macros

- #define MODEL_TYPE()
- #define MODEL_TEMPLATE_ARGS()
- #define MODEL_TEMPLATE(a, b) template MODEL_TEMPLATE_ARGS() inline a MODEL_TYPE()::b

Functions

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

8.31.1 Macro Definition Documentation

8.31.1.1 MODEL_TEMPLATE

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

8.31.1.2 MODEL_TEMPLATE_ARGS

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

8.31.1.3 MODEL_TYPE

Definition at line 91 of file model-meat.hpp.

8.31.2 Function Documentation

8.31.2.1 likelihood_()

Definition at line 56 of file model-meat.hpp.

8.31.2.2 MODEL_TEMPLATE() [1/2]

```
MODEL_TEMPLATE (

Model )
```

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

8.31.2.3 MODEL_TEMPLATE() [2/2]

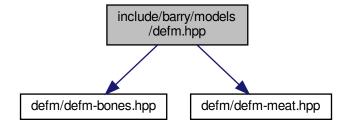
Definition at line 165 of file model-meat.hpp.

8.31.2.4 update_normalizing_constant()

Definition at line 11 of file model-meat.hpp.

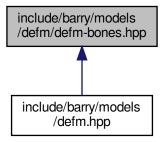
8.32 include/barry/models/defm.hpp File Reference

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



8.33 include/barry/models/defm/defm-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

- RULE_FUNCTION (rule_empty_free)
- std::vector< double > keygen_full (const phylocounters::PhyloArray &array)
- $\bullet \ \ \mathsf{bool} \ \mathsf{vec_diff} \ (\mathsf{const} \ \mathsf{std} : \mathsf{vector} < \mathsf{unsigned} \ \mathsf{int} > \&\mathsf{s}, \ \mathsf{const} \ \mathsf{std} : \mathsf{vector} < \mathsf{unsigned} \ \mathsf{int} > \&\mathsf{a})$

8.33.1 Macro Definition Documentation

8.33.1.1 INITIALIZED

```
#define INITIALIZED( )

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

Definition at line 9 of file defm-bones.hpp.

8.33.2 Function Documentation

8.33.2.1 keygen_full()

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

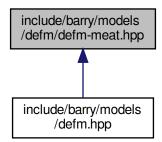
8.33.2.2 RULE_FUNCTION()

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

8.33.2.3 vec_diff()

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

8.34 include/barry/models/defm/defm-meat.hpp File Reference



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

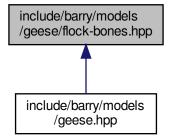
```
#include "geese/geese-node-bones.hpp"
#include "geese/geese-bones.hpp"
#include "geese/geese-meat.hpp"
#include "geese/geese-meat-constructors.hpp"
#include "geese/geese-meat-likelihood.hpp"
#include "geese/geese-meat-likelihood_exhaust.hpp"
#include "geese/geese-meat-simulate.hpp"
#include "geese/geese-meat-predict.hpp"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

Include dependency graph for geese.hpp:



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

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



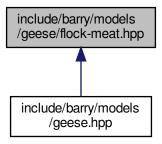
Classes

class Flock

A Flock is a group of Geese.

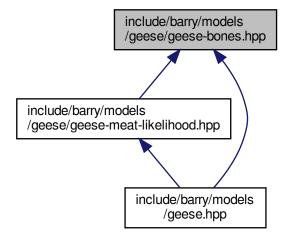
8.37 include/barry/models/geese/flock-meat.hpp File Reference

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



8.38 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)

8.38.1 Macro Definition Documentation

8.38.1.1 INITIALIZED

```
#define INITIALIZED( )

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

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

8.38.2 Function Documentation

8.38.2.1 keygen full()

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

8.38.2.2 RULE_FUNCTION()

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

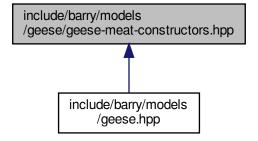
8.38.2.3 vec_diff()

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

8.38.2.4 vector_caster()

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

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



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

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



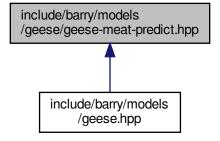


8.41 include/barry/models/geese/geese-meat-likelihood_exhaust.hpp File Reference

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

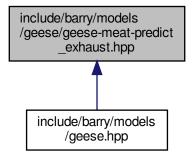


8.42 include/barry/models/geese/geese-meat-predict.hpp File Reference

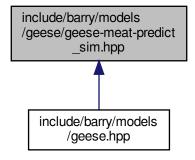


8.43 include/barry/models/geese/geese-meat-predict_exhaust.hpp File Reference

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

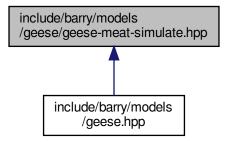


8.44 include/barry/models/geese/geese-meat-predict_sim.hpp File Reference

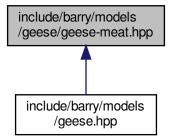


8.45 include/barry/models/geese/geese-meat-simulate.hpp File Reference

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

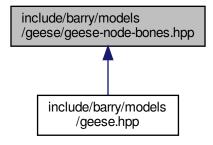


8.46 include/barry/models/geese/geese-meat.hpp File Reference



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

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



Classes

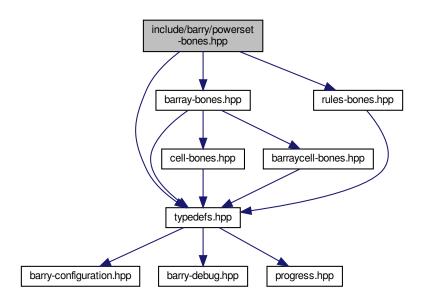
• class Node

A single node for the model.

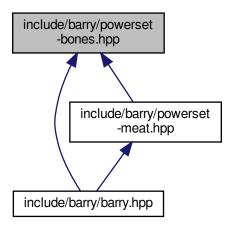
8.48 include/barry/powerset-bones.hpp File Reference

```
#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:

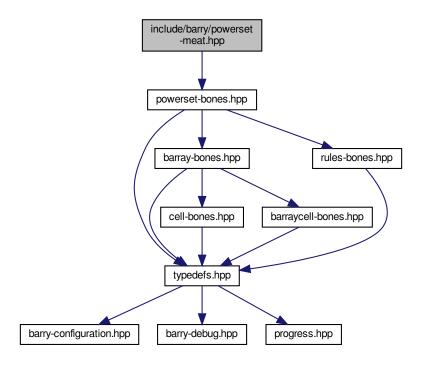


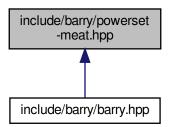
Classes

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

#include "powerset-bones.hpp"

Include dependency graph for powerset-meat.hpp:





8.50 include/barry/progress.hpp File Reference

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



Classes

class Progress
 A simple progress bar.

Macros

• #define BARRY_PROGRESS_BAR_WIDTH 80

8.50.1 Macro Definition Documentation

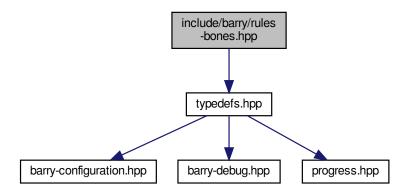
8.50.1.1 BARRY_PROGRESS_BAR_WIDTH

#define BARRY_PROGRESS_BAR_WIDTH 80

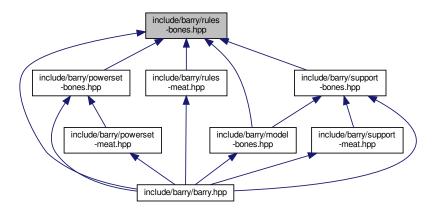
Definition at line 5 of file progress.hpp.

8.51 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)

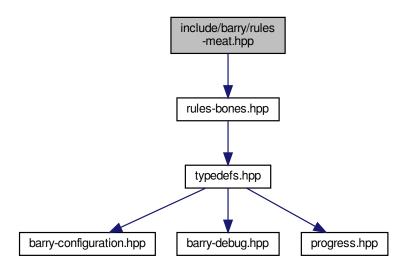
8.51.1 Function Documentation

8.51.1.1 rule_fun_default()

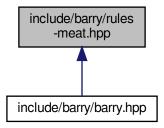
Definition at line 7 of file rules-bones.hpp.

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

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



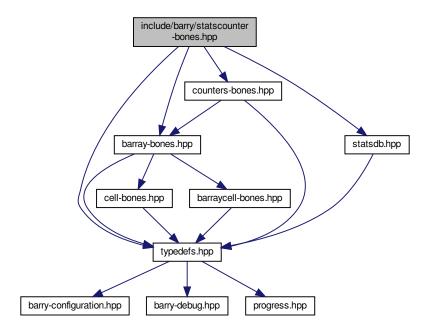
This graph shows which files directly or indirectly include this file:



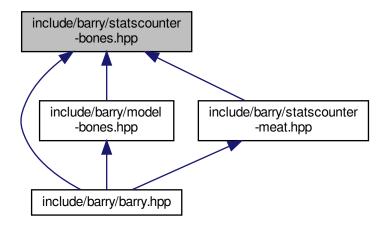
8.53 include/barry/statscounter-bones.hpp File Reference

```
#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:

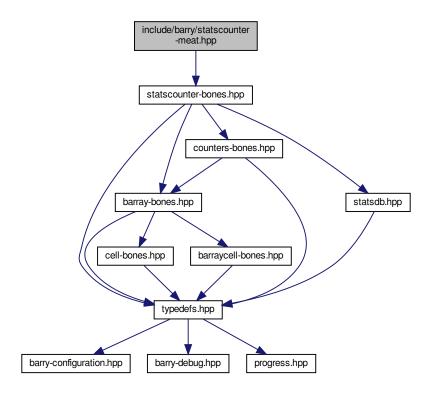


Classes

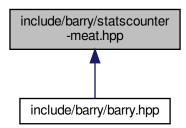
class StatsCounter < Array_Type, Data_Type >
 Count stats for a single Array.

8.54 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:



Macros

- #define STATSCOUNTER_TYPE() StatsCounter<Array_Type,Data_Type>
- #define STATSCOUNTER_TEMPLATE_ARGS() < typename Array_Type, typename Data_Type>
- #define STATSCOUNTER_TEMPLATE(a, b) template STATSCOUNTER_TEMPLATE_ARGS() inline a STATSCOUNTER_TYPE()::b

Functions

- STATSCOUNTER_TEMPLATE (, StatsCounter)(const StatsCounter< Array_Type
- EmptyArray clear ()
- STATSCOUNTER_TEMPLATE (,~StatsCounter)()
- STATSCOUNTER_TEMPLATE (void, reset_array)(const Array_Type *Array_)
- STATSCOUNTER_TEMPLATE (void, add_counter)(Counter< Array_Type
- STATSCOUNTER_TEMPLATE (void, set_counters)(Counters < Array_Type
- STATSCOUNTER_TEMPLATE (void, count_init)(uint i
- current stats resize (counters->size(), 0.0)
- for (uint n=0u;n< counters->size();++n) current stats[n]
- STATSCOUNTER_TEMPLATE (void, count_current)(uint i
- STATSCOUNTER_TEMPLATE (std::vector< std::string >, get_names)() const
- STATSCOUNTER_TEMPLATE (std::vector< std::string >, get_descriptions)() const

Variables

```
· Data_Type & counter
```

- EmptyArray = *Array
- current stats = counter.current stats
- counters = new Counters
 Array_Type,Data_Type>((*counter.counters))
- counter_deleted = false
- Data_Type * f_
- return
- Data Type * counters
- uint j

8.54.1 Macro Definition Documentation

8.54.1.1 STATSCOUNTER_TEMPLATE

Definition at line 11 of file statscounter-meat.hpp.

8.54.1.2 STATSCOUNTER TEMPLATE ARGS

```
template STATSCOUNTER_TEMPLATE_ARGS() <typename Array_Type</pre>, typename Data_Type>
```

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

8.54.1.3 STATSCOUNTER_TYPE

```
template Data_Type * STATSCOUNTER_TYPE( ) StatsCounter<Array_Type,Data_Type>
```

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

8.54.2 Function Documentation

8.54.2.1 clear()

```
EmptyArray clear ( )
```

8.54.2.2 for()

8.54.2.3 resize()

8.54.2.4 STATSCOUNTER_TEMPLATE() [1/9]

```
STATSCOUNTER_TEMPLATE (
StatsCounter ) const
```

8.54.2.5 STATSCOUNTER_TEMPLATE() [2/9]

```
STATSCOUNTER_TEMPLATE ( \sim StatsCounter )
```

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

8.54.2.6 STATSCOUNTER_TEMPLATE() [3/9]

```
STATSCOUNTER_TEMPLATE (
          std::vector< std::string > ,
          get_descriptions ) const
```

Definition at line 267 of file statscounter-meat.hpp.

8.54.2.7 STATSCOUNTER_TEMPLATE() [4/9]

Definition at line 262 of file statscounter-meat.hpp.

8.54.2.8 STATSCOUNTER_TEMPLATE() [5/9]

8.54.2.9 STATSCOUNTER_TEMPLATE() [6/9]

8.54.2.10 STATSCOUNTER_TEMPLATE() [7/9]

8.54.2.11 STATSCOUNTER_TEMPLATE() [8/9]

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

8.54.2.12 STATSCOUNTER_TEMPLATE() [9/9]

8.54.3 Variable Documentation

8.54.3.1 counter

```
Data_Type& counter

Initial value:
{
    Array = counter.Array
```

Definition at line 15 of file statscounter-meat.hpp.

8.54.3.2 counter_deleted

```
counter_deleted = false
```

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

8.54.3.3 counters

```
counters = new Counters<Array_Type, Data_Type>((*counter.counters))
```

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

8.54.3.4 counters_

```
Data_Type* counters_
Initial value:
{
    if (!counter_deleted)
```

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

8.54.3.5 current_stats

```
current_stats = counter.current_stats
```

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

8.54.3.6 EmptyArray

```
EmptyArray = *Array
```

Definition at line 20 of file statscounter-meat.hpp.

8.54.3.7 f_

```
Data_Rule_Dyn_Type f_
```

Initial value:

```
counters->add_counter(f_)
```

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

8.54.3.8 j

```
uint j
```

Initial value:

```
if (counters->size() == 0u)
    throw std::logic_error("No counters added: Cannot count without knowning what to count!")
```

Definition at line 77 of file statscounter-meat.hpp.

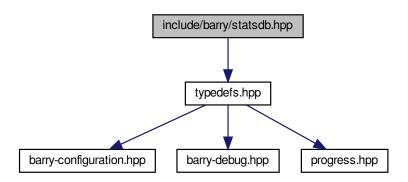
8.54.3.9 return

return

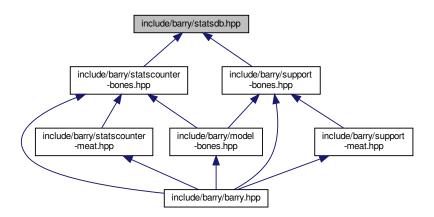
Definition at line 51 of file statscounter-meat.hpp.

8.55 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 >

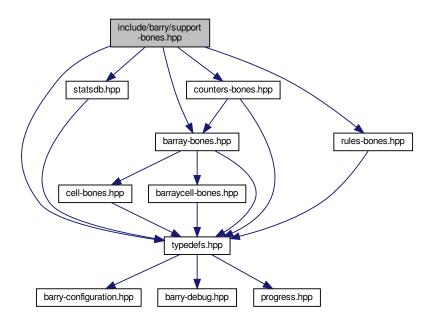
Frequency table of vectors.

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

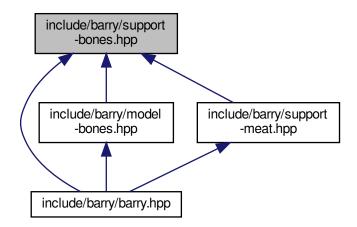
```
#include "typedefs.hpp"
#include "barray-bones.hpp"
```

```
#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:

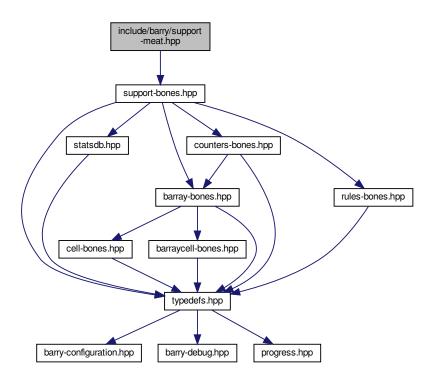


Classes

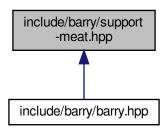
class Support < Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >
 Compute the support of sufficient statistics.

8.57 include/barry/support-meat.hpp File Reference

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



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



Macros

- #define BARRY SUPPORT MEAT HPP 1
- #define SUPPORT_TEMPLATE_ARGS()
- #define SUPPORT_TYPE()
- #define SUPPORT_TEMPLATE(a, b)

Functions

```
    SUPPORT TEMPLATE (void, init support)(std

    SUPPORT_TEMPLATE (void, reset_array)()

    SUPPORT TEMPLATE (void, reset_array)(const Array_Type &Array_)

• SUPPORT_TEMPLATE (void, calc_backend_sparse)(uint pos

    calc backend sparse (pos+1u, array bank, stats bank)

    EmptyArray insert_cell (coord_i, coord_j, EmptyArray.default_val().value, false, false)

 for (uint n=0u;n< n counters;++n)</li>

• if (rules_dyn->size() > 0u)

    if (array bank !=nullptr) array bank -> push back(EmptyArray)

if (stats_bank !=nullptr) stats_bank -> push_back(current_stats)
• EmptyArray rm_cell (coord_i, coord_j, false, false)
• if (change_stats_different > 0u)

    SUPPORT TEMPLATE (void, calc backend dense)(uint pos

• calc_backend_dense (pos+1u, array_bank, stats_bank)

    EmptyArray insert_cell (coord_i, coord_j, 1, false, false)

    SUPPORT_TEMPLATE (void, calc)(std

    SUPPORT_TEMPLATE (void, add_counter)(Counter< Array_Type</li>

• SUPPORT TEMPLATE (void, set counters)(Counters< Array Type

    SUPPORT TEMPLATE (void, add rule)(Rule < Array Type</li>

    SUPPORT_TEMPLATE (void, set_rules)(Rules< Array_Type</li>

    SUPPORT_TEMPLATE (void, add_rule_dyn)(Rule < Array_Type</li>

    SUPPORT_TEMPLATE (void, set_rules_dyn)(Rules < Array_Type</li>

    SUPPORT_TEMPLATE (bool, eval_rules_dyn)(const std

• SUPPORT_TEMPLATE (std::vector< double >, get_counts)() const

    SUPPORT TEMPLATE (std::vector< double > *, get current stats)()

    SUPPORT TEMPLATE (void, print)() const
```

Variables

```
    std::vector< Array_Type > * array_bank

    std::vector< Array Type > std::vector< std::vector< double > > * stats bank

• const size t & coord i = coordinates free[pos * 2u]

    const size t & coord j = coordinates free[pos * 2u + 1u]

    double tmp_chng

• unsigned int change_stats_different = hashes_initialized[pos] ? 0u : 1u
· & hashes [pos]

    return

    Data_Counter_Type * f_

    Data_Counter_Type * counters_

delete_counters = false
counters = counters

    Data_Rule_Type * rules_

• delete_rules = false
rules = rules
• delete rules dyn = false
• rules_dyn = rules_
```

SUPPORT_TEMPLATE (const FreqTable<> &, get_data)() const

8.57.1 Macro Definition Documentation

8.57.1.1 BARRY_SUPPORT_MEAT_HPP

```
#define BARRY_SUPPORT_MEAT_HPP 1
```

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

8.57.1.2 SUPPORT_TEMPLATE

Value:

```
template SUPPORT_TEMPLATE_ARGS() \
inline a SUPPORT_TYPE()::b
```

Definition at line 12 of file support-meat.hpp.

8.57.1.3 SUPPORT_TEMPLATE_ARGS

```
template SUPPORT_TEMPLATE_ARGS( )
```

Value:

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

8.57.1.4 SUPPORT_TYPE

```
template Data_Rule_Dyn_Type * SUPPORT_TYPE( )
```

Value:

```
Support<Array_Type,Data_Counter_Type,Data_Rule_Type,\
Data_Rule_Dyn_Type>
```

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

8.57.2 Function Documentation

8.57.2.1 calc_backend_dense()

```
calc_backend_dense (
    pos+ 1u,
    array_bank ,
    stats_bank )
```

8.57.2.2 calc_backend_sparse()

```
calc_backend_sparse (
    pos+ lu,
    array_bank ,
    stats_bank )
```

8.57.2.3 for()

```
for ( )
```

Definition at line 162 of file support-meat.hpp.

8.57.2.4 if() [1/4]

```
if (
     array_bank ! = nullptr ) -> push_back(EmptyArray)
```

8.57.2.5 if() [2/4]

```
if ( \label{eq:change_stats_different} \mbox{,} \\ \mbox{Ou })
```

Definition at line 243 of file support-meat.hpp.

8.57.2.6 if() [3/4]

```
if (
    rules_dyn-> size(),
    0u )
```

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

8.57.2.7 if() [4/4]

```
if (
    stats_bank ! = nullptr ) -> push_back(current_stats)
```

8.57.2.8 insert_cell() [1/2]

8.57.2.9 insert_cell() [2/2]

8.57.2.10 rm_cell()

8.57.2.11 SUPPORT_TEMPLATE() [1/17]

```
SUPPORT_TEMPLATE (
          bool ,
          eval_rules_dyn ) const
```

Definition at line 501 of file support-meat.hpp.

8.57.2.12 SUPPORT_TEMPLATE() [2/17]

Definition at line 550 of file support-meat.hpp.

8.57.2.13 SUPPORT_TEMPLATE() [3/17]

```
SUPPORT_TEMPLATE (
          std::vector< double > * ,
          get_current_stats )
```

Definition at line 535 of file support-meat.hpp.

8.57.2.14 SUPPORT_TEMPLATE() [4/17]

```
SUPPORT_TEMPLATE (
          std::vector< double > ,
          get_counts ) const
```

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

8.57.2.15 SUPPORT_TEMPLATE() [5/17]

8.57.2.16 SUPPORT_TEMPLATE() [6/17]

8.57.2.17 SUPPORT_TEMPLATE() [7/17]

8.57.2.18 SUPPORT_TEMPLATE() [8/17]

```
SUPPORT_TEMPLATE ( void , calc )
```

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

8.57.2.19 SUPPORT_TEMPLATE() [9/17]

8.57.2.20 SUPPORT_TEMPLATE() [10/17]

8.57.2.21 SUPPORT_TEMPLATE() [11/17]

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

8.57.2.22 SUPPORT_TEMPLATE() [12/17]

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

8.57.2.23 SUPPORT_TEMPLATE() [13/17]

```
SUPPORT_TEMPLATE (
     void ,
     reset_array )
```

Definition at line 117 of file support-meat.hpp.

8.57.2.24 SUPPORT_TEMPLATE() [14/17]

```
SUPPORT_TEMPLATE (
            void ,
            reset_array ) const &
```

Definition at line 123 of file support-meat.hpp.

8.57.2.25 SUPPORT_TEMPLATE() [15/17]

8.57.2.26 SUPPORT_TEMPLATE() [16/17]

```
SUPPORT_TEMPLATE (
     void ,
     set_rules )
```

8.57.2.27 SUPPORT_TEMPLATE() [17/17]

8.57.3 Variable Documentation

8.57.3.1 array_bank

```
std::vector< Array_Type > * array_bank
```

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

8.57.3.2 change_stats_different

```
unsigned int change_stats_different = hashes_initialized[pos] ? Ou : 1u
```

Definition at line 161 of file support-meat.hpp.

8.57.3.3 coord_i

```
const size_t & coord_i = coordinates_free[pos * 2u]
```

Definition at line 147 of file support-meat.hpp.

8.57.3.4 coord_j

```
const size_t & coord_j = coordinates_free[pos * 2u + 1u]
```

Definition at line 148 of file support-meat.hpp.

8.57.3.5 counters

```
counters = counters_
```

Definition at line 429 of file support-meat.hpp.

8.57.3.6 counters_

```
Data_Counter_Type* counters_
Initial value:
{
    if (delete_counters)
```

Definition at line 422 of file support-meat.hpp.

8.57.3.7 delete_counters

```
delete_counters = false
```

Definition at line 428 of file support-meat.hpp.

8.57.3.8 delete_rules

```
delete_rules = false
```

Definition at line 462 of file support-meat.hpp.

8.57.3.9 delete_rules_dyn

```
delete_rules_dyn = false
```

Definition at line 494 of file support-meat.hpp.

8.57.3.10 else

Definition at line 216 of file support-meat.hpp.

8.57.3.11 f_

```
Data_Rule_Dyn_Type f_
Initial value:
{
    counters->add_counter(f_)
```

Definition at line 404 of file support-meat.hpp.

8.57.3.12 hashes

```
& hashes
```

Definition at line 221 of file support-meat.hpp.

8.57.3.13 return

return

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

8.57.3.14 rules

```
rules = rules_
```

Definition at line 463 of file support-meat.hpp.

8.57.3.15 rules_

```
Data_Rule_Dyn_Type * rules_
Initial value:
{
    if (delete_rules)
```

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

8.57.3.16 rules_dyn

```
rules_dyn = rules_
```

Definition at line 495 of file support-meat.hpp.

8.57.3.17 stats_bank

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

Initial value:
{
    if (pos >= coordiantes_n_free)
        return
```

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

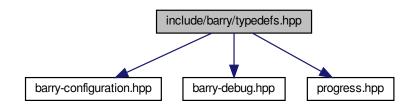
8.57.3.18 tmp_chng

```
double tmp_chng
```

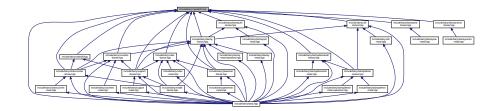
Definition at line 160 of file support-meat.hpp.

8.58 include/barry/typedefs.hpp File Reference

```
#include "barry-configuration.hpp"
#include "barry-debug.hpp"
#include "progress.hpp"
Include dependency graph for typedefs.hpp:
```



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



Classes

```
    class Entries < Cell_Type >
        A wrapper class to store source, target, val from a BArray object.

    struct vecHasher < T >
```

Namespaces

CHECK

Integer constants used to specify which cell should be check.

• EXISTS

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

Typedefs

```
· typedef unsigned int uint
```

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

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

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

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

```
    template < typename Array_Type, typename Data_Type >
    using Counter_fun_type = std::function < double(const Array_Type &, uint, uint, Data_Type &) >
        Counter and rule functions.
```

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

Functions

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

```
    template<typename T >
        bool vec_equal (const std::vector< T > &a, const std::vector< T > &b)
    Compares if -a- and -b- are equal.
```

```
    template<typename T >
        bool vec_equal_approx (const std::vector< T > &a, const std::vector< T > &b, double eps=1e-100)
```

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

8.58.1 Typedef Documentation

8.58.1.1 Col_type

```
template<typename Cell_Type >
using Col_type = Map< uint, Cell<Cell_Type>* >
```

Definition at line 71 of file typedefs.hpp.

8.58.1.2 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 148 of file typedefs.hpp.

8.58.1.3 Counts_type

```
typedef std::vector< std::pair< std::vector<double>, uint > > Counts_type
```

Definition at line 52 of file typedefs.hpp.

8.58.1.4 MapVec_type

```
template<typename Ta = double, typename Tb = uint>
using MapVec_type = std::unordered_map< std::vector< Ta >, Tb, vecHasher<Ta> >
```

Definition at line 129 of file typedefs.hpp.

8.58.1.5 Row_type

```
template<typename Cell_Type >
using Row_type = Map< uint, Cell<Cell_Type> >
```

Definition at line 68 of file typedefs.hpp.

8.58.1.6 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 151 of file typedefs.hpp.

8.58.1.7 uint

```
typedef unsigned int uint
```

Definition at line 18 of file typedefs.hpp.

8.58.2 Function Documentation

8.58.2.1 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 162 of file typedefs.hpp.

8.58.2.2 vec_equal_approx()

Definition at line 180 of file typedefs.hpp.

8.58.2.3 vec inner_prod() [1/2]

Definition at line 223 of file typedefs.hpp.

8.58.2.4 vec_inner_prod() [2/2]

Definition at line 200 of file typedefs.hpp.

8.59 README.md File Reference

Index

```
\simBArray
                                                             Progress, 177
                                                        \simRule
     BArray< Cell Type, Data Type >, 37
\simBArrayCell
                                                             Rule < Array_Type, Data_Type >, 179
     BArrayCell< Cell_Type, Data_Type >, 48
                                                        \simRules
~BArrayCell const
                                                             Rules < Array_Type, Data_Type >, 181
     BArrayCell_const< Cell_Type, Data_Type >, 50
                                                        \simStatsCounter
{\sim}\mathsf{BArrayDense}
                                                             StatsCounter < Array_Type, Data_Type >, 185
     BArrayDense < Cell_Type, Data_Type >, 56
                                                        \simSupport
                                                                          Array_Type,
                                                                                          Data Counter Type,
\simBArrayDenseCell
                                                             Support<
     BArrayDenseCell< Cell_Type, Data_Type >, 68
                                                                  Data_Rule_Type, Data_Rule_Dyn_Type >,
\simBArrayRow
                                                                  190
     BArrayRow< Cell Type, Data Type >, 81
                                                        active
~BArrayRow const
                                                             Cell< Cell_Type >, 96
     BArrayRow_const< Cell_Type, Data_Type >, 83
                                                        add
\simBArrayVector
                                                             barray-meat.hpp, 218
     BArrayVector< Cell Type, Data Type >, 86
                                                             barraydense-meat.hpp, 245
~BArrayVector const
                                                             Cell< Cell_Type >, 94, 95
     BArrayVector_const< Cell_Type, Data_Type >, 89
                                                             FreqTable < T >, 119
\simCell
                                                        add array
     Cell < Cell Type >, 93
                                                             Model<
                                                                          Array_Type,
                                                                                           Data_Counter_Type,
\simConstBArrayRowIter
                                                                  Data_Rule_Type, Data_Rule_Dyn_Type >,
     ConstBArrayRowlter< Cell Type, Data Type >, 98
                                                                  144
\simCounter
                                                        add counter
     Counter< Array Type, Data Type >, 101
                                                             Counters < Array_Type, Data_Type >, 106
\simCounters
                                                             Model<
                                                                          Array Type,
                                                                                          Data Counter Type,
    Counters < Array_Type, Data_Type >, 105
                                                                  Data_Rule_Type, Data_Rule_Dyn_Type >,
\simEntries
                                                                  144, 145
    Entries < Cell_Type >, 109
                                                             StatsCounter< Array_Type, Data_Type >, 185
\simFlock
                                                             Support<
                                                                          Array_Type,
                                                                                          Data_Counter_Type,
     Flock, 112
                                                                  Data_Rule_Type, Data_Rule_Dyn_Type >,
\simFreqTable
                                                                  191
     FreqTable < T >, 118
                                                        add_data
\simGeese
                                                             Flock, 112
    Geese, 125, 126
                                                        add rule
\simModel
                                                             Model<
                                                                          Array_Type,
                                                                                          Data_Counter_Type,
                                  Data Counter Type,
     Model<
                 Array_Type,
                                                                  Data_Rule_Type, Data_Rule_Dyn_Type >,
         \label{eq:def:def:Data_Rule_Dyn_Type} \ \ \mathsf{Data} \_ \mathsf{Rule} \_ \mathsf{Dyn} \_ \mathsf{Type} \ \ >,
         144
                                                             PowerSet< Array_Type, Data_Rule_Type >, 172,
\simNetCounterData
                                                                  173
    NetCounterData, 155
                                                             Rules < Array_Type, Data_Type >, 182
\simNetworkData
                                                             Support<
                                                                          Array Type,
                                                                                          Data Counter Type,
    NetworkData, 158
                                                                  {\tt Data\_Rule\_Type}, \quad {\tt Data\_Rule\_Dyn\_Type} \quad >, \quad
\simNode
    Node, 160
                                                        add rule dyn
\simPhyloRuleDynData
                                                             Model<
                                                                                          Data Counter Type,
                                                                          Array Type,
     PhyloRuleDynData, 169
                                                                  Data_Rule_Type, Data_Rule_Dyn_Type >,
\simPowerSet
                                                                  145
     PowerSet < Array_Type, Data_Rule_Type >, 172
                                                                                           Data_Counter_Type,
                                                             Support<
                                                                           Array_Type,
\simProgress
                                                                  Data_Rule_Type, Data_Rule_Dyn_Type >,
```

191	row, 44
annotations	set_data, 44
Node, 161	swap_cells, 45
ans	swap_cols, 45
barray-meat.hpp, 209, 218	swap_rows, 45
barraydense-meat.hpp, 235, 246	toggle_cell, 45
Array	toggle_lock, 45
ConstBArrayRowIter< Cell_Type, Data_Type >, 98	transpose, 46
array	visited, 47
Node, 161	zero_col, 46
Array_	zero_row, 46
barray-meat.hpp, 218	barray-bones.hpp
array_bank	BARRAY_BONES_HPP, 200
support-meat.hpp, 341	barray-meat-operators.hpp
arrays	BARRAY_TEMPLATE, 202-204 BARRAY_TEMPLATE_ARGS, 202, 204
Node, 162 AS_ONE	BARRAY_TYPE, 202, 204
	BARRY BARRAY MEAT OPERATORS HPP,
EXISTS, 31 as_vector	202
FreqTable < T >, 119	COL, 203
AS_ZERO	for, 204
EXISTS, 31	operator(), 205
at	rhs, 205
PhyloCounterData, 166	ROW, 203
Thylodounior Bata, 100	this, 205
BArray	barray-meat.hpp
BArray< Cell_Type, Data_Type >, 36, 37	add, 218
BArray< Cell_Type, Data_Type >, 33	ans, 209, 218
\sim BArray, 37	Array_, 218
BArray, 36, 37	BARRAY_TEMPLATE, 208–213
BArrayCell< Cell_Type, Data_Type >, 46	BARRAY_TEMPLATE_ARGS, 208
BArrayCell_const< Cell_Type, Data_Type >, 46	BARRAY_TYPE, 208
clear, 37	check_bounds, 218
col, 37	check_exists, 218
D, 38	COL, 209, 213
default_val, 38	col0, 219
flush_data, 38	const, 219
get_cell, 38	copy_data, 219
get_col_vec, 38, 39	data, 219
get_entries, 39	delete_data, 219
get_row_vec, 39	delete_data_, 220
insert_cell, 39, 40	else, 220
is_dense, 40	false, 220
is_empty, 40	first, 220
ncol, 40	for, 213, 214
nnozero, 41	i1, 220
nrow, 41	if, 214–216
operator*=, 41	j, 221
operator(), 41	j0, 221
operator+=, 41, 42	j1, 221
operator-=, 42	M, 217, 221
operator/=, 42	M_, 221
operator=, 43 operator==, 43	N, 222
out_of_range, 43	NCells, 222
print, 43	report, 222
reserve, 43	resize, 217
resize, 44	return, 217, 222
rm_cell, 44	ROW, 209, 217
III_00II, TT	

row0, 222	col, 56, 57
search, 223	colsum, 57
source, 223	D, 57
target, 223	default_val, 57
v, 223	get_cell, 57
value, 223	get_col_vec, 58
BARRAY_BONES_HPP	get_data, 58
barray-bones.hpp, 200	get_entries, 58
BARRAY_TEMPLATE	get_row_vec, 58, 59
barray-meat-operators.hpp, 202-204	insert_cell, 59
barray-meat.hpp, 208-213	is_dense, 59
BARRAY_TEMPLATE_ARGS	is_empty, 59
barray-meat-operators.hpp, 202, 204	ncol, 60
barray-meat.hpp, 208	nnozero, 60
BARRAY_TYPE	nrow, 60
barray-meat-operators.hpp, 202, 204	operator*=, 60
barray-meat.hpp, 208	operator(), 60
BArrayCell	operator+=, 61
BArrayCell< Cell_Type, Data_Type >, 48	operator-=, 61
BArrayCell< Cell Type, Data Type >, 47	operator/=, 62
~BArrayCell, 48	operator=, 62
BArray Cell_Type, Data_Type >, 46	operator==, 62
BArrayCell, 48	out_of_range, 62
	-
operator Cell_Type, 48	print, 62
operator*=, 48	reserve, 63
operator+=, 48	resize, 63
operator-=, 49	rm_cell, 63
operator/=, 49	row, 63
operator=, 49	rowsum, 63
operator==, 49	set_data, 64
BArrayCell_const	swap_cells, 64
BArrayCell_const< Cell_Type, Data_Type >, 50	swap_cols, 64
BArrayCell_const< Cell_Type, Data_Type >, 50	swap_rows, 64
\sim BArrayCell_const, 50	toggle_cell, 65
BArray< Cell_Type, Data_Type >, 46	toggle_lock, 65
BArrayCell_const, 50	transpose, 65
operator Cell_Type, 51	visited, 67
operator!=, 51	zero_col, 65
operator<, 51	zero_row, 65
operator<=, 51	barraydense-bones.hpp
operator>, 51	BARRY_BARRAYDENSE_BONES_HPP, 227
operator>=, 52	barraydense-meat-operators.hpp
operator==, 51	BARRY_BARRAYDENSE_MEAT_OPERATORS_HPP,
BArrayDense	228
BArrayDense < Cell_Type, Data_Type >, 55, 56	BDENSE_TEMPLATE, 228-230
BArrayDense< Cell_Type, Data_Type >, 52	BDENSE_TEMPLATE_ARGS, 228, 230
∼BArrayDense, 56	BDENSE_TYPE, 228, 230
BArrayDense, 55, 56	COL, 229
BArrayDenseCell< Cell_Type, Data_Type >, 66,	POS, 229
70	POS_N, 229
BArrayDenseCol < Cell_Type, Data_Type >, 66, 73	ROW, 229
BArrayDenseCol_const< Cell_Type, Data_Type >,	barraydense-meat.hpp
66	add, 245
BArrayDenseRow< Cell_Type, Data_Type >, 66,	ans, 235, 246
77	BDENSE_TEMPLATE, 233, 235–242
BArrayDenseRow_const< Cell_Type, Data_Type	BDENSE_TEMPLATE_ARGS, 233
>, 66	BDENSE_TYPE, 234
clear, 56	check_bounds, 246
51041, 00	Shook_boahab, £70

check exists, 246	operator==, 69
	•
COL, 234	barraydensecell-bones.hpp
col, 246	POS, 252
const, 247	barraydensecell-meat.hpp
copy_data, 247	POS, 254
data, 247	BArrayDenseCell_const< Cell_Type, Data_Type >, 71
delete_data, 247	BArrayDenseCol < Cell_Type, Data_Type >, 73
delete_data_, 247	BArrayDenseCol_const< Cell_Type, Data_Type >
el, 248	75
el_colsums, 248	BArrayDenseRow< Cell_Type, Data_Type >, 77
el rowsums, 248	
-	BArrayDenseRow_const< Cell_Type, Data_Type
else, 248	>, 80
false, 248	BArrayDenseCol
for, 242	BArrayDenseCol< Cell_Type, Data_Type >, 71
i1, 249	BArrayDenseCol< Cell_Type, Data_Type >, 71
if, 242	BArrayDense< Cell_Type, Data_Type >, 66, 73
insert_cell, 243	BArrayDenseCell< Cell_Type, Data_Type >, 70
j, 249	73
j0, 249	BArrayDenseCell const< Cell Type, Data Type
j1, 249	>, 73
M, 243, 249	BArrayDenseCol, 71
M_, 249	begin, 72
N, 250	end, 72
POS, 234	operator(), 72
POS_N, 234	size, 72
report, 250	barraydensecol-bones.hpp
resize, 243, 244	POS, 255
return, 250	POS_N, 255
rm_cell, 244, 245	ZERO_CELL, 255
ROW, 234	BArrayDenseCol_const
source, 250	BArrayDenseCol_const< Cell_Type, Data_Type >
target, 250	74
v, 251	BArrayDenseCol_const< Cell_Type, Data_Type >, 73
	BArrayDense< Cell_Type, Data_Type >, 66
va_end, 245	
va_start, 245	BArrayDenseCell< Cell_Type, Data_Type >, 70,
val0, 251	75
val1, 251	BArrayDenseCell_const< Cell_Type, Data_Type
value, 251	>, 75
vprintf, 245	BArrayDenseCol_const, 74
ZERO_CELL, 235	begin, 74
BArrayDenseCell	end, 74
BArrayDenseCell< Cell_Type, Data_Type >, 68	operator(), 74
BArrayDenseCell< Cell_Type, Data_Type >, 67	size, 75
~BArrayDenseCell, 68	BArrayDenseRow
BArrayDense< Cell_Type, Data_Type >, 66, 70	BArrayDenseRow< Cell_Type, Data_Type >, 76
BArrayDenseCell, 68	BArrayDenseRow< Cell_Type, Data_Type >, 75
BArrayDenseCol Cell_Type, Data_Type >, 70, 73	BArrayDense Cell_Type, Data_Type >, 66, 77
BArrayDenseCol_const< Cell_Type, Data_Type >,	BArrayDenseCell Cell_Type, Data_Type >, 77
70, 75	BArrayDenseCell_const< Cell_Type, Data_Type
BArrayDenseRow< Cell_Type, Data_Type >, 77	>, 77
BArrayDenseRow_const< Cell_Type, Data_Type	BArrayDenseRow, 76
>, 80	begin, 76
operator Cell_Type, 68	end, 76
operator*=, 68	operator(), 77
operator+=, 69	size, 77
operator-=, 69	barraydenserow-bones.hpp
operator/=, 69	POS, 256
operator=, 69	POS_N, 256
υροιαισι-, σο	1 OO_11, 200

7FD0_0FU_0F0	amayatay 00
ZERO_CELL, 256	operator=, 88
BArrayDenseRow_const	operator==, 88
BArrayDenseRow_const< Cell_Type, Data_Type	size, 88
>, 78	barrayvector-meat.hpp
BArrayDenseRow_const< Cell_Type, Data_Type >, 78	BARRY_BARRAYVECTOR_MEAT_HPP, 262
BArrayDense < Cell_Type, Data_Type >, 66	BArrayVector_const
BArrayDenseCell< Cell_Type, Data_Type >, 80	BArrayVector_const< Cell_Type, Data_Type >, 89
BArrayDenseCell_const< Cell_Type, Data_Type	BArrayVector_const< Cell_Type, Data_Type >, 88
>, 80	~BArrayVector_const, 89
BArrayDenseRow_const, 78	BArrayVector const, 89
begin, 79	begin, 90
end, 79	end, 90
operator(), 79	
	is_col, 90
size, 79	is_row, 90
BArrayRow	operator std::vector< Cell_Type >, 90
BArrayRow< Cell_Type, Data_Type >, 81	operator!=, 90
BArrayRow< Cell_Type, Data_Type >, 80	operator<, 91
\sim BArrayRow, 81	operator<=, 91
BArrayRow, 81	operator>, 91
operator BArrayRow< Cell_Type, Data_Type >, 81	operator>=, 91
operator*=, 81	operator==, 91
operator+=, 81	size, 92
operator-=, 81	barry, 29
operator/=, 82	barry-configuration.hpp
operator=, 82	BARRY_CHECK_SUPPORT, 263
operator==, 82	BARRY_ISFINITE, 263
barrayrow-meat.hpp	BARRY_MAX_NUM_ELEMENTS, 263
BARRY_BARRAYROW_MEAT_HPP, 258	BARRY_SAFE_EXP, 263
BROW_TEMPLATE, ABOX, 850	Map, 263
BROW_TEMPLATE_ARGS, 259	printf_barry, 263
BROW_TYPE, 259	barry-debug.hpp
BArrayRow_const	BARRY_DEBUG_LEVEL, 264
BArrayRow_const< Cell_Type, Data_Type >, 83	barry-macros.hpp
BArrayRow_const< Cell_Type, Data_Type >, 82	BARRY_ONE, 265
\sim BArrayRow_const, 83	BARRY_ONE_DENSE, 265
BArrayRow_const, 83	BARRY_UNUSED, 265
operator BArrayRow_const< Cell_Type, Data_Type	BARRY_ZERO, 265
>, 83	BARRY_ZERO_DENSE, 265
operator!=, 83	barry.hpp
operator<, 83	BARRY_HPP, 267
operator<=, 84	BARRY VERSION, 267
operator>, 84	COUNTER FUNCTION, 267
operator>=, 84	COUNTER LAMBDA, 268
operator==, 84	RULE FUNCTION, 268
BArrayVector	RULE_LAMBDA, 268
BArrayVector< Cell_Type, Data_Type >, 85	barry::counters, 29
BArrayVector Cell_Type, Data_Type >, 84	barry::counters; 25
~BArrayVector, 86	-
•	barry::counters::phylo, 30
BArrayVector, 85	BARRY_BARRAY_MEAT_OPERATORS_HPP
begin, 86	barray-meat-operators.hpp, 202
end, 86	BARRY_BARRAYDENSE_BONES_HPP
is_col, 86	barraydense-bones.hpp, 227
is_row, 86	BARRY_BARRAYDENSE_MEAT_OPERATORS_HPP
operator std::vector< Cell_Type >, 87	barraydense-meat-operators.hpp, 228
operator*=, 87	BARRY_BARRAYROW_MEAT_HPP
operator+=, 87	barrayrow-meat.hpp, 258
operator-=, 87	BARRY_BARRAYVECTOR_MEAT_HPP
operator/=, 87	barrayvector-meat.hpp, 262

BARRY_CHECK_SUPPORT	barrayrow-meat.hpp, 258–260
barry-configuration.hpp, 263 BARRY DEBUG LEVEL	BROW_TEMPLATE_ARGS barrayrow-meat.hpp, 259
barry-debug.hpp, 264	BROW TYPE
BARRY HPP	barrayrow-meat.hpp, 259
barry.hpp, 267	•
BARRY_ISFINITE	calc
barry-configuration.hpp, 263	PowerSet < Array_Type, Data_Rule_Type >, 173
BARRY_MAX_NUM_ELEMENTS	Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,
barry-configuration.hpp, 263	192
BARRY_ONE	calc_backend_dense
barry-macros.hpp, 265 BARRY_ONE_DENSE	support-meat.hpp, 336
barry-macros.hpp, 265	calc_backend_sparse
BARRY_PROGRESS_BAR_WIDTH	support-meat.hpp, 337
progress.hpp, 322	calc_reduced_sequence
BARRY_SAFE_EXP	Geese, 126
barry-configuration.hpp, 263	calc_sequence
BARRY_SUPPORT_MEAT_HPP	Geese, 126, 127
support-meat.hpp, 335	Cell
BARRY_UNUSED	Cell < Cell Type >, 93, 94
barry-macros.hpp, 265	Cell< Cell_Type $>$, 92 \sim Cell, 93
BARRY_VERSION	active, 96
barry.hpp, 267	add, 94, 95
BARRY_ZERO barry-macros.hpp, 265	Cell, 93, 94
BARRY_ZERO_DENSE	operator Cell_Type, 95
barry-macros.hpp, 265	operator!=, 95
BARRY_ZERO_NETWORK	operator=, 95, 96
network.hpp, 292	operator==, 96
BARRY_ZERO_NETWORK_DENSE	value, 96
network.hpp, 293	visited, 96
BDENSE_TEMPLATE	Cell_const< Cell_Type >, 97
barraydense-meat-operators.hpp, 228-230	change_stats
barraydense-meat.hpp, 233, 235–242	Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,
BDENSE_TEMPLATE_ARGS	195
barraydense-meat-operators.hpp, 228, 230	change_stats_different
barraydense-meat.hpp, 233 BDENSE_TYPE	support-meat.hpp, 342
barraydense-meat-operators.hpp, 228, 230	CHECK, 30
barraydense-meat.hpp, 234	BOTH, 30
begin	NONE, 30
BArrayDenseCol< Cell_Type, Data_Type >, 72	ONE, 30
${\sf BArrayDenseCol_const} {< Cell_Type, Data_Type >},$	TWO, 30
74	check_bounds
BArrayDenseRow< Cell_Type, Data_Type >, 76	barray-meat.hpp, 218 barraydense-meat.hpp, 246
BArrayDenseRow_const< Cell_Type, Data_Type	check_exists
>, 79	barray-meat.hpp, 218
BArrayVector< Cell_Type, Data_Type >, 86 BArrayVector_const< Cell_Type, Data_Type >, 90	barraydense-meat.hpp, 246
PhyloCounterData, 166	clear
PowerSet < Array_Type, Data_Rule_Type >, 173	BArray< Cell_Type, Data_Type >, 37
blengths	BArrayDense < Cell_Type, Data_Type >, 56
NodeData, 165	FreqTable < T >, 119
BOTH	statscounter-meat.hpp, 328
CHECK, 30	COL
EXISTS, 31	barray-meat-operators.hpp, 203 barray-meat.hpp, 209, 213
BROW_TEMPLATE	barraydense-meat-operators.hpp, 229
	- may solve most operatorompp,

	barraydense-meat.hpp, 234	count
col		Counter< Array_Type, Data_Type >, 102
	BArray< Cell_Type, Data_Type >, 37	count_all
	BArrayDense < Cell_Type, Data_Type >, 56, 57	StatsCounter< Array_Type, Data_Type >, 186
10	barraydense-meat.hpp, 246	count_current
col0		StatsCounter < Array_Type, Data_Type >, 186
Cal	barray-meat.hpp, 219	count_fun Counter< Array Type Data Type > 103
COI_	_type typedefs.hpp, 347	Counter< Array_Type, Data_Type >, 103 counters-meat.hpp, 275
coln	ames	count_fun_
COIII	Flock, 112	counters-meat.hpp, 279
	Geese, 127	count_init
	Model< Array_Type, Data_Counter_Type,	StatsCounter< Array_Type, Data_Type >, 186
	Data_Rule_Type, Data_Rule_Dyn_Type >,	Counter
	146	Counter< Array_Type, Data_Type >, 100, 101
cols	um	counter
	BArrayDense < Cell_Type, Data_Type >, 57	counters-meat.hpp, 279
cond	ditional_prob	statscounter-meat.hpp, 330
	Model < Array_Type, Data_Counter_Type,	Counter< Array_Type, Data_Type >, 99
	Data_Rule_Type, Data_Rule_Dyn_Type >,	\sim Counter, 101
	146	count, 102
cons	st	count_fun, 103
	barray-meat.hpp, 219	Counter, 100, 101
	barraydense-meat.hpp, 247	data, 103
Con	stBArrayRowIter	desc, 103
_	ConstBArrayRowlter< Cell_Type, Data_Type >, 98	get_description, 102
Con	stBArrayRowlter< Cell_Type, Data_Type >, 97	get_name, 102
	~ConstBArrayRowlter, 98	init, 102
	Array, 98	init_fun, 103
	ConstBArrayRowlter, 98	name, 103
	current_col, 99	operator=, 102
	current_row, 99 iter, 99	counter_
coor		counters-meat.hpp, 279 counter_absdiff
COOI	support-meat.hpp, 342	Network counters, 13
coor		counter_co_opt
000.	support-meat.hpp, 342	Phylo counters, 21
coor	diantes_n_free	counter_cogain
	Support< Array_Type, Data_Counter_Type,	Phylo counters, 21
	Data_Rule_Type, Data_Rule_Dyn_Type >,	counter css census01
	195	network-css.hpp, 285
coor	diantes_n_locked	counter_css_census02
	Support< Array_Type, Data_Counter_Type,	network-css.hpp, 286
	Data_Rule_Type, Data_Rule_Dyn_Type >,	counter_css_census03
	195	network-css.hpp, 286
coor	dinates_free	counter_css_census04
	PowerSet < Array_Type, Data_Rule_Type >, 175	network-css.hpp, 286
	Support< Array_Type, Data_Counter_Type,	counter_css_census05
	Data_Rule_Type, Data_Rule_Dyn_Type >,	network-css.hpp, 286
	195	counter_css_census06
coor	dinates_locked	network-css.hpp, 287
	PowerSet < Array_Type, Data_Rule_Type >, 175	counter_css_census07
	Support Array_Type, Data_Counter_Type,	network-css.hpp, 287
	Data_Rule_Type, Data_Rule_Dyn_Type >,	counter_css_census08
00:0:	196	network-css.hpp, 287
copy	/_data	counter_css_census09
	barray-meat.hpp, 219 barraydense-meat.hpp, 247	network-css.hpp, 287 counter_css_census10
	Darrayaction inicatifipp, <u>LT</u>	00011101_000_001100010

network-css.hpp, 288	Phylo counters, 23
counter_css_completely_false_recip_comiss	counter_neofun_a2b
network-css.hpp, 288	Phylo counters, 24
counter_css_completely_false_recip_omiss	counter_nodecov
network-css.hpp, 288	Network counters, 16
counter_css_mixed_recip	counter_nodeicov
network-css.hpp, 288	Network counters, 17
counter_css_partially_false_recip_commi	counter_nodematch
network-css.hpp, 289	Network counters, 17
counter_css_partially_false_recip_omiss	counter_nodeocov
network-css.hpp, 289	Network counters, 17
counter_ctriads	counter odegree
Network counters, 13, 14	Network counters, 17
counter_degree	counter_odegree15
Network counters, 14	Network counters, 18
counter_deleted	counter_ostar2
statscounter-meat.hpp, 330	Network counters, 18
counter_density	counter_overall_changes
Network counters, 14	Phylo counters, 24
counter diff	counter_overall_gains
-	Phylo counters, 24
Network counters, 14	· · · · · · · · · · · · · · · · · · ·
counter_edges	counter_overall_gains_from_0
Network counters, 14	Phylo counters, 24
Counter_fun_type	counter_overall_loss
typedefs.hpp, 347	Phylo counters, 25
COUNTER_FUNCTION	counter_pairwise_first_gain
barry.hpp, 267	Phylo counters, 25
counter_gains	counter_pairwise_neofun_singlefun
Phylo counters, 21	Phylo counters, 25
counter_gains_from_0	counter_pairwise_overall_change
Phylo counters, 21	Phylo counters, 25
counter_gains_k_offspring	counter_pairwise_preserving
Phylo counters, 22	Phylo counters, 26
counter_genes_changing	counter_preserve_pseudogene
Phylo counters, 22	Phylo counters, 26
counter_idegree	counter_prop_genes_changing
Network counters, 15	Phylo counters, 26
counter_idegree15	counter_subfun
Network counters, 15	Phylo counters, 26
counter_isolates	COUNTER_TEMPLATE
Network counters, 15, 16	counters-meat.hpp, 273, 275, 276
counter_istar2	COUNTER_TEMPLATE_ARGS
Network counters, 16	counters-meat.hpp, 274
counter_k_genes_changing	counter_ttriads
Phylo counters, 22	Network counters, 18, 19
COUNTER_LAMBDA	COUNTER_TYPE
barry.hpp, 268	counters-meat.hpp, 274
counter_less_than_p_prop_genes_changing	Counters
Phylo counters, 22	Counters < Array_Type, Data_Type >, 105
counter_longest	counters-meat.hpp, 276
Phylo counters, 23	counters
counter_loss	statscounter-meat.hpp, 330
Phylo counters, 23	support-meat.hpp, 342
counter_maxfuns	Counters< Array_Type, Data_Type >, 104
Phylo counters, 23	~Counters, 105
counter_mutual	add_counter, 106
Network counters, 16	Counters, 105
counter_neofun	get_descriptions, 106
Joannol_HodiuH	got_accomptions, roo

get_names, 106	network-css.hpp, 285
operator=, 106, 107	CSS_SIZE
operator[], 107	network-css.hpp, 285
size, 107	CSS_TRUE_CELLS
counters-meat.hpp	network-css.hpp, 285
count_fun, 275	current_col
count_fun_, 279	ConstBArrayRowIter< Cell_Type, Data_Type >, 99
counter, 279	current_row
counter_, 279	ConstBArrayRowIter< Cell_Type, Data_Type >, 99
COUNTER_TEMPLATE, 273, 275, 276	current_stats
COUNTER_TEMPLATE_ARGS, 274	statscounter-meat.hpp, 330
COUNTER_TYPE, 274	Support< Array_Type, Data_Counter_Type,
COUNTERS TEMPLATE 274 276 277	Data_Rule_Type, Data_Rule_Dyn_Type >,
COUNTERS_TEMPLATE, 274, 276, 277	196
COUNTERS_TEMPLATE_ARGS, 274 COUNTERS TYPE, 274	D
data, 278	BArray< Cell Type, Data Type >, 38
data_, 279	BArrayDense< Cell_Type, Data_Type >, 57
desc, 278	Rule < Array_Type, Data_Type >, 180
desc_, 280	dat
i, 280	Flock, 116
init_fun, 278	data
init_fun_, 280	barray-meat.hpp, 219
j, 280	barraydense-meat.hpp, 247
name, 278	Counter< Array_Type, Data_Type >, 103
name_, 280	counters-meat.hpp, 278
noexcept, 281	PowerSet < Array_Type, Data_Rule_Type >, 175
return, 281	data_
this, 281	counters-meat.hpp, 279
counters_	DEFAULT_DUPLICATION
statscounter-meat.hpp, 330	phylo.hpp, 299
support-meat.hpp, 342	default_val
COUNTERS_TEMPLATE	BArray Danas Call Type, Data Type >, 38
counters-meat.hpp, 274, 276, 277	BArrayDense < Cell_Type, Data_Type >, 57 defm-bones.hpp
COUNTERS_TEMPLATE_ARGS	INITIALIZED, 309
counters-meat.hpp, 274	keygen_full, 310
COUNTERS_TYPE	RULE_FUNCTION, 310
counters-meat.hpp, 274 Counting, 11	vec_diff, 310
counts	delete counters
PhyloRuleDynData, 169	Support< Array_Type, Data_Counter_Type,
Counts_type	Data_Rule_Type, Data_Rule_Dyn_Type >,
typedefs.hpp, 347	196
CSS_APPEND	support-meat.hpp, 342
network-css.hpp, 283	delete_data
CSS_CASE_ELSE	barray-meat.hpp, 219
network-css.hpp, 283	barraydense-meat.hpp, 247
CSS_CASE_PERCEIVED	delete_data_
network-css.hpp, 284	barray-meat.hpp, 220
CSS_CASE_TRUTH	barraydense-meat.hpp, 247
network-css.hpp, 284	delete_rengine
CSS_CHECK_SIZE	Geese, 138
network-css.hpp, 284	delete_rules
CSS_CHECK_SIZE_INIT	Support< Array_Type, Data_Counter_Type,
network-css.hpp, 284	Data_Rule_Type, Data_Rule_Dyn_Type >,
CSS_NET_COUNTER_LAMBDA_INIT	196 support-meat hop, 343
network-css.hpp, 284	support-meat.hpp, 343 delete_rules_dyn
CSS PERCEIVED CELLS	doloto_raios_ayri

Support< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 196	val, 110 eval_rules_dyn Support< Array_Type, Data_Counter_Type.
support-meat.hpp, 343 delete_support	Data_Rule_Type, Data_Rule_Dyn_Type >
Geese, 139	EXISTS, 31
desc	AS_ONE, 31
Counter< Array_Type, Data_Type >, 103	AS_ZERO, 31
counters-meat.hpp, 278	BOTH, 31
desc_	NONE, 32
counters-meat.hpp, 280	ONE, 32
directed	TWO, 32
NetworkData, 158	UKNOWN, 32
DUPL_DUPL	,
phylo.hpp, 299	f_
DUPL_EITH	statscounter-meat.hpp, 331
phylo.hpp, 299	support-meat.hpp, 343
DUPL_SPEC	false
phylo.hpp, 299	barray-meat.hpp, 220
duplication	barraydense-meat.hpp, 248
Node, 162	first
NodeData, 165	barray-meat.hpp, 220
PhyloRuleDynData, 169	Flock, 110
	~Flock, 112
el	add_data, 112
barraydense-meat.hpp, 248	colnames, 112
el_colsums	dat, 116
barraydense-meat.hpp, 248	Flock, 112
el_rowsums	get_counters, 113
barraydense-meat.hpp, 248	get_model, 113
else	get_stats_support, 113
barray-meat.hpp, 220	get_stats_target, 113
barraydense-meat.hpp, 248	get_support_fun, 113
support-meat.hpp, 343	init, 113
empty PhyloCounterDate 166	initialized, 116
PhyloCounterData, 166 EmptyArray	likelihood_joint, 114 model, 117
• • •	nfunctions, 117
PowerSet< Array_Type, Data_Rule_Type >, 175	nfuns, 114
statscounter-meat.hpp, 331 end	nleafs, 114
BArrayDenseCol< Cell_Type, Data_Type >, 72	nnodes, 114
BArrayDenseCol_const < Cell_Type, Data_Type >, 72	nterms, 115
74	ntrees, 115
BArrayDenseRow< Cell Type, Data Type >, 76	operator(), 115
BArrayDenseRow_const< Cell_Type, Data_Type	parse_polytomies, 115
>, 79	print, 116
BArrayVector< Cell_Type, Data_Type >, 86	rengine, 117
BArrayVector_const< Cell_Type, Data_Type >, 90	set_seed, 116
PhyloCounterData, 167	support_size, 116
PowerSet< Array_Type, Data_Rule_Type >, 173	flush_data
Progress, 178	BArray< Cell_Type, Data_Type >, 38
Entries	for
Entries < Cell_Type >, 109	barray-meat-operators.hpp, 204
Entries Cell_Type >, 108	barray-meat-operators.npp, 204 barray-meat.hpp, 213, 214
~Entries, 109	barraydense-meat.hpp, 242
Entries, 109	statscounter-meat.hpp, 328
resize, 109	support-meat.hpp, 337
source, 109	FreqTable
target, 110	FreqTable < T >, 118

FreqTable 118	geese-bones.hpp
∼FreqTable, 118	INITIALIZED, 313
add, 119	keygen_full, 313 RULE_FUNCTION, 313
as_vector, 119 clear, 119	vec_diff, 313
FreqTable, 118	vector_caster, 314
get_data, 119 get_index, 119	gen_key Model< Array Type, Data Counter Type,
make_hash, 120	3= 31 / = = 31 /
print, 120	Data_Rule_Type, Data_Rule_Dyn_Type >, 146
•	
reserve, 120	get_annotated_nodes Geese, 127
size, 120	
Geese, 121	get_arrays2support Model< Array_Type, Data_Counter_Type,
∼Geese, 125, 126	Data_Rule_Type, Data_Rule_Dyn_Type >,
calc_reduced_sequence, 126	147
calc_sequence, 126, 127	
colnames, 127	get_cell RArray < Coll Type Data Type > 38
delete_rengine, 138	BArray< Cell_Type, Data_Type >, 38 BArrayDense< Cell Type, Data Type >, 57
delete_support, 139	
Geese, 124–126	get_col_vec
get_annotated_nodes, 127	BArray Cell_Type, Data_Type >, 38, 39
get_counters, 127, 128	BArrayDense < Cell_Type, Data_Type >, 58
get_model, 128	get_counters
get_probabilities, 128	Flock, 113
get_rengine, 128	Geese, 127, 128
get_states, 129	Model < Array_Type, Data_Counter_Type,
get_support_fun, 129	Data_Rule_Type, Data_Rule_Dyn_Type >,
inherit_support, 129, 130	147
init, 130	PhyloCounterData, 167
init_node, 130	StatsCounter< Array_Type, Data_Type >, 186
initialized, 139	Support< Array_Type, Data_Counter_Type,
likelihood, 130, 131	Data_Rule_Type, Data_Rule_Dyn_Type >,
likelihood_exhaust, 131	192
map_to_nodes, 139	get_counts
nannotations, 131	Support< Array_Type, Data_Counter_Type,
nfunctions, 139	Data_Rule_Type, Data_Rule_Dyn_Type >,
nfuns, 132	192
nleafs, 132	get_current_stats
nnodes, 132	Support< Array_Type, Data_Counter_Type,
nodes, 139	Data_Rule_Type, Data_Rule_Dyn_Type >,
nterms, 133	193
observed_counts, 133	get_data
operator=, 133, 134	BArrayDense < Cell_Type, Data_Type >, 58
parse_polytomies, 134	FreqTable < T >, 119
predict, 134	PowerSet < Array_Type, Data_Rule_Type >, 173
predict_backend, 135	Support< Array_Type, Data_Counter_Type,
predict exhaust, 135	Data_Rule_Type, Data_Rule_Dyn_Type >,
predict_exhaust_backend, 135, 136	193
predict_sim, 136	get_data_ptr
print, 136	PowerSet < Array_Type, Data_Rule_Type >, 174
print_observed_counts, 137	get_description
pset_loc, 139	Counter< Array_Type, Data_Type >, 102
reduced_sequence, 140	get_descriptions
sequence, 140	Counters < Array_Type, Data_Type >, 106
set_seed, 137	StatsCounter< Array_Type, Data_Type >, 186
simulate, 137	get_entries
	BArray< Cell_Type, Data_Type >, 39
support_size, 138 update_annotations, 138	BArrayDense< Cell_Type, Data_Type >, 58
apaate annotations, 100	

get_index	get_seq
FreqTable < T >, 119	Rules< Array_Type, Data_Type >, 182
get_last_name	get_states
phylo.hpp, 304	Geese, 129
get_model	get_stats_support
Flock, 113	Flock, 113
Geese, 128	Model< Array_Type, Data_Counter_Type,
get_name	Data_Rule_Type, Data_Rule_Dyn_Type >,
Counter< Array_Type, Data_Type >, 102	149
get_names	get_stats_target
Counters< Array_Type, Data_Type >, 106	Flock, 113
StatsCounter< Array_Type, Data_Type >, 186	Model< Array_Type, Data_Counter_Type,
get_norm_const	$Data_Rule_Type, Data_Rule_Dyn_Type >,$
Model< Array_Type, Data_Counter_Type,	149
Data_Rule_Type, Data_Rule_Dyn_Type >,	get_support_fun
147	Flock, 113
get_parent	Geese, 129
Node, 161	Model < Array_Type, Data_Counter_Type,
get_probabilities	Data_Rule_Type, Data_Rule_Dyn_Type >,
Geese, 128	149
get_pset	
Model < Array_Type, Data_Counter_Type,	hashes
Data_Rule_Type, Data_Rule_Dyn_Type >,	Support< Array_Type, Data_Counter_Type,
147	Data_Rule_Type, Data_Rule_Dyn_Type >,
get_pset_arrays	197
Model< Array_Type, Data_Counter_Type,	support-meat.hpp, 343
Data_Rule_Type, Data_Rule_Dyn_Type >,	hashes_initialized
147	Support< Array_Type, Data_Counter_Type,
get_pset_probs	Data_Rule_Type, Data_Rule_Dyn_Type >,
Model< Array_Type, Data_Counter_Type,	197
Data_Rule_Type, Data_Rule_Dyn_Type >,	
148	i
get_pset_stats	counters-meat.hpp, 280
Model< Array_Type, Data_Counter_Type,	i1
Data_Rule_Type, Data_Rule_Dyn_Type >,	barray-meat.hpp, 220
148	barraydense-meat.hpp, 249
get_rengine	id
Geese, 128	Node, 162
Model< Array_Type, Data_Counter_Type,	if
Data_Rule_Type, Data_Rule_Dyn_Type >,	barray-meat.hpp, 214–216
148	barraydense-meat.hpp, 242
get_row_vec	support-meat.hpp, 337
BArray< Cell_Type, Data_Type >, 39	IF_MATCHES
BArrayDense< Cell Type, Data Type >, 58, 59	phylo.hpp, 300
get_rules	IF_NOTMATCHES
Model< Array_Type, Data_Counter_Type,	phylo.hpp, 300
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/barray-bones.hpp, 199
148	include/barry/barray-iterator.hpp, 200
Support< Array_Type, Data_Counter_Type,	include/barry/barray-meat-operators.hpp, 201
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/barray-meat.hpp, 206
193	include/barry/barraycell-bones.hpp, 224
get_rules_dyn	include/barry/barraycell-meat.hpp, 224
Model< Array_Type, Data_Counter_Type,	include/barry/barraydense-bones.hpp, 225
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/barraydense-meat-operators.hpp, 227
149	include/barry/barraydense-meat.hpp, 231
	include/barry/barraydensecell-bones.hpp, 252
Support< Array_Type, Data_Counter_Type,	include/barry/barraydensecell-meat.hpp, 253
Data_Rule_Type, Data_Rule_Dyn_Type >,	include/barry/barraydensecol-bones.hpp, 254
193	include/barry/barraydenserow-bones.hpp, 255

include/barry/barrayrow-bones.hpp, 257	Counter< Array_Type, Data_Type >, 102
include/barry/barrayrow-meat.hpp, 257	Flock, 113
include/barry/barrayvector-bones.hpp, 260	Geese, 130
include/barry/barrayvector-meat.hpp, 261	init_fun
include/barry/barry-configuration.hpp, 262	Counter< Array_Type, Data_Type >, 103
include/barry/barry-debug.hpp, 264	counters-meat.hpp, 278
include/barry/barry-macros.hpp, 264	init_fun_
include/barry/barry.hpp, 266	counters-meat.hpp, 280
include/barry/cell-bones.hpp, 269	init_node
include/barry/cell-meat.hpp, 269	Geese, 130
include/barry/col-bones.hpp, 270	init_support
include/barry/counters-bones.hpp, 270	PowerSet < Array_Type, Data_Rule_Type >, 174
include/barry/counters-meat.hpp, 272	Support< Array_Type, Data_Counter_Type,
include/barry/counters/network-css.hpp, 282	Data_Rule_Type, Data_Rule_Dyn_Type >,
include/barry/counters/network.hpp, 290	193
include/barry/counters/phylo.hpp, 297	INITIALIZED
include/barry/model-bones.hpp, 304	defm-bones.hpp, 309
include/barry/model-meat.hpp, 306	geese-bones.hpp, 313
include/barry/models/defm.hpp, 308	initialized
include/barry/models/defm/defm-bones.hpp, 309	Flock, 116
include/barry/models/defm/defm-meat.hpp, 310	Geese, 139
include/barry/models/geese.hpp, 311	insert_cell
include/barry/models/geese/flock-bones.hpp, 311	BArray< Cell_Type, Data_Type >, 39, 40
include/barry/models/geese/flock-meat.hpp, 312	BArrayDense< Cell_Type, Data_Type >, 59
include/barry/models/geese/geese-bones.hpp, 312	barraydense-meat.hpp, 243
include/barry/models/geese/geese-meat-constructors.hpp	o, support-meat.hpp, 338
314	is_col
include/barry/models/geese/geese-meat-likelihood.hpp,	BArrayVector< Cell_Type, Data_Type >, 86
315	BArrayVector_const< Cell_Type, Data_Type >, 90
include/barry/models/geese/geese-meat-likelihood_exhau	usis <u>h</u> pp ņse
316	BArray< Cell_Type, Data_Type >, 40
include/barry/models/geese/geese-meat-predict.hpp,	BArrayDense< Cell_Type, Data_Type >, 59
316	IS_DUPLICATION
include/barry/models/geese/geese-meat-predict_exhaust.	
317	IS_EITHER
include/barry/models/geese/geese-meat-predict_sim.hpp	, phylo.hpp, 300
317	is_empty
include/barry/models/geese/geese-meat-simulate.hpp,	BArray< Cell_Type, Data_Type >, 40
318	
310	BArrayDense < Cell_Type, Data_Type >, 59
include/barry/models/geese/geese-meat.hpp, 318	
	BArrayDense< Cell_Type, Data_Type >, 59
include/barry/models/geese/geese-meat.hpp, 318	BArrayDense< Cell_Type, Data_Type >, 59 is_leaf
include/barry/models/geese/geese-meat.hpp, 318 include/barry/models/geese/geese-node-bones.hpp,	BArrayDense< Cell_Type, Data_Type >, 59 is_leaf Node, 161
include/barry/models/geese/geese-meat.hpp, 318 include/barry/models/geese/geese-node-bones.hpp, 319	BArrayDense< Cell_Type, Data_Type >, 59 is_leaf Node, 161 is_row
include/barry/models/geese/geese-meat.hpp, 318 include/barry/models/geese/geese-node-bones.hpp, 319 include/barry/powerset-bones.hpp, 319	BArrayDense< Cell_Type, Data_Type >, 59 is_leaf Node, 161 is_row BArrayVector< Cell_Type, Data_Type >, 86
include/barry/models/geese/geese-meat.hpp, 318 include/barry/models/geese/geese-node-bones.hpp, 319 include/barry/powerset-bones.hpp, 319 include/barry/powerset-meat.hpp, 320	BArrayDense< Cell_Type, Data_Type >, 59 is_leaf Node, 161 is_row BArrayVector< Cell_Type, Data_Type >, 86 BArrayVector_const< Cell_Type, Data_Type >, 90
include/barry/models/geese/geese-meat.hpp, 318 include/barry/models/geese/geese-node-bones.hpp, 319 include/barry/powerset-bones.hpp, 319 include/barry/powerset-meat.hpp, 320 include/barry/progress.hpp, 322 include/barry/rules-bones.hpp, 322 include/barry/rules-meat.hpp, 324	BArrayDense< Cell_Type, Data_Type >, 59 is_leaf Node, 161 is_row BArrayVector< Cell_Type, Data_Type >, 86 BArrayVector_const< Cell_Type, Data_Type >, 90 IS_SPECIATION phylo.hpp, 300 iter
include/barry/models/geese/geese-meat.hpp, 318 include/barry/models/geese/geese-node-bones.hpp, 319 include/barry/powerset-bones.hpp, 319 include/barry/powerset-meat.hpp, 320 include/barry/progress.hpp, 322 include/barry/rules-bones.hpp, 322	BArrayDense< Cell_Type, Data_Type >, 59 is_leaf Node, 161 is_row BArrayVector< Cell_Type, Data_Type >, 86 BArrayVector_const< Cell_Type, Data_Type >, 90 IS_SPECIATION phylo.hpp, 300
include/barry/models/geese/geese-meat.hpp, 318 include/barry/models/geese/geese-node-bones.hpp, 319 include/barry/powerset-bones.hpp, 319 include/barry/powerset-meat.hpp, 320 include/barry/rogress.hpp, 322 include/barry/rules-bones.hpp, 322 include/barry/rules-meat.hpp, 324 include/barry/statscounter-bones.hpp, 324 include/barry/statscounter-meat.hpp, 326	BArrayDense< Cell_Type, Data_Type >, 59 is_leaf Node, 161 is_row BArrayVector< Cell_Type, Data_Type >, 86 BArrayVector_const< Cell_Type, Data_Type >, 90 IS_SPECIATION phylo.hpp, 300 iter ConstBArrayRowIter< Cell_Type, Data_Type >, 99
include/barry/models/geese/geese-meat.hpp, 318 include/barry/models/geese/geese-node-bones.hpp, 319 include/barry/powerset-bones.hpp, 319 include/barry/powerset-meat.hpp, 320 include/barry/progress.hpp, 322 include/barry/rules-bones.hpp, 322 include/barry/rules-meat.hpp, 324 include/barry/statscounter-bones.hpp, 324 include/barry/statscounter-meat.hpp, 326 include/barry/statsdb.hpp, 332	BArrayDense < Cell_Type, Data_Type >, 59 is_leaf Node, 161 is_row BArrayVector < Cell_Type, Data_Type >, 86 BArrayVector_const < Cell_Type, Data_Type >, 90 IS_SPECIATION phylo.hpp, 300 iter ConstBArrayRowIter < Cell_Type, Data_Type >, 99 j
include/barry/models/geese/geese-meat.hpp, 318 include/barry/models/geese/geese-node-bones.hpp, 319 include/barry/powerset-bones.hpp, 319 include/barry/powerset-meat.hpp, 320 include/barry/progress.hpp, 322 include/barry/rules-bones.hpp, 322 include/barry/rules-meat.hpp, 324 include/barry/statscounter-bones.hpp, 324 include/barry/statscounter-meat.hpp, 326 include/barry/statsdb.hpp, 332 include/barry/support-bones.hpp, 332	BArrayDense < Cell_Type, Data_Type >, 59 is_leaf Node, 161 is_row BArrayVector < Cell_Type, Data_Type >, 86 BArrayVector_const < Cell_Type, Data_Type >, 90 IS_SPECIATION phylo.hpp, 300 iter ConstBArrayRowIter < Cell_Type, Data_Type >, 99 j barray-meat.hpp, 221
include/barry/models/geese/geese-meat.hpp, 318 include/barry/models/geese/geese-node-bones.hpp, 319 include/barry/powerset-bones.hpp, 319 include/barry/powerset-meat.hpp, 320 include/barry/progress.hpp, 322 include/barry/rules-bones.hpp, 322 include/barry/rules-meat.hpp, 324 include/barry/statscounter-bones.hpp, 324 include/barry/statscounter-meat.hpp, 326 include/barry/statsdb.hpp, 332 include/barry/support-bones.hpp, 332 include/barry/support-meat.hpp, 334	BArrayDense< Cell_Type, Data_Type >, 59 is_leaf
include/barry/models/geese/geese-meat.hpp, 318 include/barry/models/geese/geese-node-bones.hpp, 319 include/barry/powerset-bones.hpp, 319 include/barry/powerset-meat.hpp, 320 include/barry/progress.hpp, 322 include/barry/rules-bones.hpp, 322 include/barry/rules-meat.hpp, 324 include/barry/statscounter-bones.hpp, 324 include/barry/statscounter-meat.hpp, 326 include/barry/statsdb.hpp, 332 include/barry/support-bones.hpp, 332 include/barry/support-meat.hpp, 334 include/barry/typedefs.hpp, 345	BArrayDense< Cell_Type, Data_Type >, 59 is_leaf
include/barry/models/geese/geese-meat.hpp, 318 include/barry/models/geese/geese-node-bones.hpp, 319 include/barry/powerset-bones.hpp, 319 include/barry/powerset-meat.hpp, 320 include/barry/rogress.hpp, 322 include/barry/rules-bones.hpp, 322 include/barry/rules-meat.hpp, 324 include/barry/statscounter-bones.hpp, 324 include/barry/statscounter-meat.hpp, 326 include/barry/statsdb.hpp, 332 include/barry/support-bones.hpp, 332 include/barry/support-meat.hpp, 334 include/barry/typedefs.hpp, 345 indices	BArrayDense < Cell_Type, Data_Type >, 59 is_leaf Node, 161 is_row BArrayVector < Cell_Type, Data_Type >, 86 BArrayVector_const < Cell_Type, Data_Type >, 90 IS_SPECIATION phylo.hpp, 300 iter ConstBArrayRowlter < Cell_Type, Data_Type >, 99 j barray-meat.hpp, 221 barraydense-meat.hpp, 249 counters-meat.hpp, 280 statscounter-meat.hpp, 331
include/barry/models/geese/geese-meat.hpp, 318 include/barry/models/geese/geese-node-bones.hpp, 319 include/barry/powerset-bones.hpp, 319 include/barry/powerset-meat.hpp, 320 include/barry/rogress.hpp, 322 include/barry/rules-bones.hpp, 322 include/barry/rules-meat.hpp, 324 include/barry/statscounter-bones.hpp, 324 include/barry/statscounter-meat.hpp, 326 include/barry/statsdb.hpp, 332 include/barry/support-bones.hpp, 332 include/barry/support-meat.hpp, 334 include/barry/typedefs.hpp, 345 indices NetCounterData, 156	BArrayDense < Cell_Type, Data_Type >, 59 is_leaf Node, 161 is_row BArrayVector < Cell_Type, Data_Type >, 86 BArrayVector_const < Cell_Type, Data_Type >, 90 IS_SPECIATION phylo.hpp, 300 iter ConstBArrayRowlter < Cell_Type, Data_Type >, 99 j barray-meat.hpp, 221 barraydense-meat.hpp, 249 counters-meat.hpp, 280 statscounter-meat.hpp, 331 j0
include/barry/models/geese/geese-meat.hpp, 318 include/barry/models/geese/geese-node-bones.hpp, 319 include/barry/powerset-bones.hpp, 319 include/barry/powerset-meat.hpp, 320 include/barry/progress.hpp, 322 include/barry/rules-bones.hpp, 322 include/barry/rules-meat.hpp, 324 include/barry/statscounter-bones.hpp, 324 include/barry/statscounter-meat.hpp, 326 include/barry/statsdb.hpp, 332 include/barry/support-bones.hpp, 332 include/barry/support-meat.hpp, 334 include/barry/typedefs.hpp, 345 indices NetCounterData, 156 inherit_support	BArrayDense < Cell_Type, Data_Type >, 59 is_leaf Node, 161 is_row BArrayVector < Cell_Type, Data_Type >, 86 BArrayVector_const < Cell_Type, Data_Type >, 90 IS_SPECIATION phylo.hpp, 300 iter ConstBArrayRowlter < Cell_Type, Data_Type >, 99 j barray-meat.hpp, 221 barraydense-meat.hpp, 249 counters-meat.hpp, 280 statscounter-meat.hpp, 331 j0 barray-meat.hpp, 221
include/barry/models/geese/geese-meat.hpp, 318 include/barry/models/geese/geese-node-bones.hpp, 319 include/barry/powerset-bones.hpp, 319 include/barry/powerset-meat.hpp, 320 include/barry/rogress.hpp, 322 include/barry/rules-bones.hpp, 322 include/barry/rules-meat.hpp, 324 include/barry/statscounter-bones.hpp, 324 include/barry/statscounter-meat.hpp, 326 include/barry/statsdb.hpp, 332 include/barry/support-bones.hpp, 332 include/barry/support-meat.hpp, 334 include/barry/typedefs.hpp, 345 indices NetCounterData, 156	BArrayDense < Cell_Type, Data_Type >, 59 is_leaf Node, 161 is_row BArrayVector < Cell_Type, Data_Type >, 86 BArrayVector_const < Cell_Type, Data_Type >, 90 IS_SPECIATION phylo.hpp, 300 iter ConstBArrayRowlter < Cell_Type, Data_Type >, 99 j barray-meat.hpp, 221 barraydense-meat.hpp, 249 counters-meat.hpp, 280 statscounter-meat.hpp, 331 j0 barray-meat.hpp, 221 barraydense-meat.hpp, 249
include/barry/models/geese/geese-meat.hpp, 318 include/barry/models/geese/geese-node-bones.hpp, 319 include/barry/powerset-bones.hpp, 319 include/barry/powerset-meat.hpp, 320 include/barry/progress.hpp, 322 include/barry/rules-bones.hpp, 322 include/barry/rules-meat.hpp, 324 include/barry/statscounter-bones.hpp, 324 include/barry/statscounter-meat.hpp, 326 include/barry/statsdb.hpp, 332 include/barry/support-bones.hpp, 332 include/barry/support-meat.hpp, 334 include/barry/typedefs.hpp, 345 indices NetCounterData, 156 inherit_support	BArrayDense < Cell_Type, Data_Type >, 59 is_leaf Node, 161 is_row BArrayVector < Cell_Type, Data_Type >, 86 BArrayVector_const < Cell_Type, Data_Type >, 90 IS_SPECIATION phylo.hpp, 300 iter ConstBArrayRowlter < Cell_Type, Data_Type >, 99 j barray-meat.hpp, 221 barraydense-meat.hpp, 249 counters-meat.hpp, 280 statscounter-meat.hpp, 331 j0 barray-meat.hpp, 221

barraydense-meat.hpp, 249	add_array, 144
1.6.16	add_counter, 144, 145
keygen_default	add_rule, 145
model-bones.hpp, 305	add_rule_dyn, 145
keygen_full	colnames, 146
defm-bones.hpp, 310	conditional_prob, 146
geese-bones.hpp, 313	gen_key, 146
lb	get_arrays2support, 147
PhyloRuleDynData, 169	get_counters, 147
likelihood	get_norm_const, 147
Geese, 130, 131	get_pset, 147
Model < Array_Type, Data_Counter_Type,	get_pset_arrays, 147
Data_Rule_Type, Data_Rule_Dyn_Type >,	get_pset_probs, 148
149, 150	get_pset_stats, 148
likelihood	get_rengine, 148
model-meat.hpp, 307	get_rules, 148
likelihood_exhaust	get_rules_dyn, 149
Geese, 131	get_stats_support, 149
likelihood joint	get_stats_target, 149
Flock, 114	get_support_fun, 149
likelihood total	likelihood, 149, 150
Model < Array_Type, Data_Counter_Type,	likelihood_total, 150
Data_Rule_Type, Data_Rule_Dyn_Type >,	Model, 143
150	nterms, 150
	operator=, 151
M	print, 151
barray-meat.hpp, 217, 221	print_stats, 151
barraydense-meat.hpp, 243, 249	sample, 151
PowerSet< Array_Type, Data_Rule_Type >, 175	set_counters, 152
Support< Array_Type, Data_Counter_Type,	set_keygen, 152
Data_Rule_Type, Data_Rule_Dyn_Type >,	set_rengine, 152
197	set_rules, 152
M	set_rules_dyn, 152
barray-meat.hpp, 221	set_seed, 153
barraydense-meat.hpp, 249	set_transform_model, 153
MAKE DUPL VARS	size, 153
phylo.hpp, 301	size_unique, 153
make_hash	store_psets, 154
FreqTable $<$ T $>$, 120	support_size, 154
Мар	transform_model, 154
barry-configuration.hpp, 263	model-bones.hpp
map_to_nodes	keygen_default, 305
Geese, 139	model-meat.hpp
MapVec_type	likelihood_, 307
typedefs.hpp, 348	MODEL_TEMPLATE, ADCC, 200
max_num_elements	MODEL_TEMPLATE_ARGS, 306
Support< Array_Type, Data_Counter_Type,	MODEL_TYPE, 307
Data_Rule_Type, Data_Rule_Dyn_Type >,	update_normalizing_constant, 308
197	MODEL_TEMPLATE
Model	model-meat.hpp, 306, 307
Model< Array_Type, Data_Counter_Type,	MODEL_TEMPLATE_ARGS
Data_Rule_Type, Data_Rule_Dyn_Type >,	model-meat.hpp, 306
143	MODEL_TYPE
model	model-meat.hpp, 307
Flock, 117	N
Model < Array_Type, Data_Counter_Type, Data_Rule_Type	
Data_Rule_Dyn_Type >, 140	barraydense-meat.hpp, 250
\sim Model, 144	PowerSet < Array Type, Data Rule Type >. 176

Support< Array_Type, Data_Counter_Type,	counter_idegree15, 15
Data_Rule_Type, Data_Rule_Dyn_Type >,	counter_isolates, 15, 16
197	counter_istar2, 16
n_counters	counter_mutual, 16
Support< Array_Type, Data_Counter_Type,	counter_nodecov, 16
Data_Rule_Type, Data_Rule_Dyn_Type >,	counter_nodeicov, 17
198	counter_nodematch, 17
n_free	counter_nodeocov, 17
PowerSet < Array_Type, Data_Rule_Type >, 176	counter_odegree, 17
n_locked	counter_odegree15, 18
PowerSet < Array_Type, Data_Rule_Type >, 176	counter_ostar2, 18
name	counter_ttriads, 18, 19
Counter< Array_Type, Data_Type >, 103	NETWORK_COUNTER, 19
counters-meat.hpp, 278	network-css.hpp
name_	counter_css_census01, 285
counters-meat.hpp, 280	counter_css_census02, 286
nannotations	counter_css_census03, 286
Geese, 131	counter_css_census04, 286
narray	counter_css_census05, 286
Node, 162	counter_css_census06, 287
NCells	counter_css_census07, 287
barray-meat.hpp, 222	counter_css_census08, 287
ncol	counter_css_census09, 287
BArray< Cell_Type, Data_Type >, 40	counter_css_census10, 288
BArrayDense < Cell_Type, Data_Type >, 60	counter_css_completely_false_recip_comiss, 288
NET_C_DATA_IDX	counter_css_completely_false_recip_omiss, 288
network.hpp, 293	counter_css_mixed_recip, 288
NET_C_DATA_NUM	counter_css_partially_false_recip_commi, 289
network.hpp, 293	counter_css_partially_false_recip_omiss, 289
NetCounter	CSS_APPEND, 283
network.hpp, 295	CSS_CASE_ELSE, 283
NetCounterData, 154	CSS_CASE_PERCEIVED, 284
∼NetCounterData, 155	CSS_CASE_TRUTH, 284
indices, 156	CSS_CHECK_SIZE, 284
NetCounterData, 155	CSS_CHECK_SIZE_INIT, 284
numbers, 156	CSS_NET_COUNTER_LAMBDA_INIT, 284
NetCounters	CSS_PERCEIVED_CELLS, 285
network.hpp, 295	CSS_SIZE, 285
NetModel	CSS_TRUE_CELLS, 285
network.hpp, 295	network.hpp
NetRule	BARRY_ZERO_NETWORK, 292
network.hpp, 295	BARRY_ZERO_NETWORK_DENSE, 293
NetRules	NET_C_DATA_IDX, 293
network.hpp, 295	NET_C_DATA_NUM, 293
NetStatsCounter	NetCounter, 295
network.hpp, 295	NetCounters, 295
NetSupport	NetModel, 295
network.hpp, 296	NetRule, 295
Network	NetRules, 295
network.hpp, 296	NetStatsCounter, 295
Network counters, 12	NetSupport, 296
counter_absdiff, 13	Network, 296
counter_ctriads, 13, 14	NETWORK_COUNTER, 293
counter_degree, 14	NETWORK_COUNTER_LAMBDA, 293
counter_density, 14	NETWORK RULE, 294
counter_diff, 14	NETWORK_RULE_LAMBDA, 294
counter_edges, 14	NetworkDense, 296
counter_idegree, 15	NETWORKDENSE_COUNTER_LAMBDA, 294

rules_zerodiag, 296	states, 165
NETWORK_COUNTER	nodes
Network counters, 19	Geese, 139
network.hpp, 293	noexcept
NETWORK_COUNTER_LAMBDA	counters-meat.hpp, 281
network.hpp, 293	noffspring
NETWORK_RULE	Node, 161
network.hpp, 294	NONE
NETWORK_RULE_LAMBDA	CHECK, 30
network.hpp, 294	EXISTS, 32
NetworkData, 156	nrow
~NetworkData, 158	BArray< Cell_Type, Data_Type >, 41
directed, 158	BArrayDense < Cell_Type, Data_Type >, 60
NetworkData, 157	nterms
vertex_attr, 158	Flock, 115
NetworkDense	Geese, 133
network.hpp, 296	Model< Array_Type, Data_Counter_Type
NETWORKDENSE_COUNTER_LAMBDA	Data_Rule_Type, Data_Rule_Dyn_Type >
network.hpp, 294	150
next	ntrees
Progress, 178	Flock, 115
nfunctions	numbers
Flock, 117	NetCounterData, 156
Geese, 139	abserved counts
nfuns	observed_counts Geese, 133
Flock, 114	offspring
Geese, 132	Node, 162
nleafs	ONE
Flock, 114	CHECK, 30
Geese, 132	EXISTS, 32
nnodes	operator BArrayRow< Cell_Type, Data_Type >
Flock, 114	BArrayRow< Cell_Type, Data_Type >, 81
Geese, 132	operator BArrayRow_const< Cell_Type, Data_Type >
nnozero	BArrayRow_const< Cell_Type, Data_Type >, 83
BArray < Cell_Type, Data_Type >, 41	operator Cell_Type
BArrayDense < Cell_Type, Data_Type >, 60	BArrayCell< Cell_Type, Data_Type >, 48
Node, 158	BArrayCell_const< Cell_Type, Data_Type >, 51
~Node, 160	BArrayDenseCell< Cell_Type, Data_Type >, 68
annotations, 161	Cell< Cell_Type >, 95
array, 161	operator std::vector< Cell Type >
arrays, 162	BArrayVector< Cell_Type, Data_Type >, 87
duplication, 162	BArrayVector const< Cell Type, Data Type >, 90
get_parent, 161	operator!=
id, 162	BArrayCell_const< Cell_Type, Data_Type >, 51
is_leaf, 161 narray, 162	BArrayRow_const< Cell_Type, Data_Type >, 83
	BArrayVector_const< Cell_Type, Data_Type >, 90
Node, 159, 160 noffspring, 161	Cell< Cell_Type >, 95
offspring, 162	operator<
ord, 163	BArrayCell_const< Cell_Type, Data_Type >, 51
parent, 163	BArrayRow_const< Cell_Type, Data_Type >, 83
parent, 163 probability, 163	BArrayVector_const< Cell_Type, Data_Type >, 91
subtree_prob, 163	operator<=
-	BArrayCell_const< Cell_Type, Data_Type >, 51
visited, 163	BArrayRow_const< Cell_Type, Data_Type >, 84
NodeData, 164	BArrayVector_const< Cell_Type, Data_Type >, 91
blengths, 165	operator>
duplication, 165 NodeData, 164	BArrayCell_const< Cell_Type, Data_Type >, 51
Nouchata, 104	BArrayRow_const< Cell_Type, Data_Type >, 84

BArrayVector_const< Cell_Type, Data_Type >, 91 operator>= BArrayCell_const< Cell_Type, Data_Type >, 52 BArrayRow_const< Cell_Type, Data_Type >, 84	Geese, 133, 134 Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >, 151
BArrayVector_const< Cell_Type, Data_Type >, 91	Rules< Array_Type, Data_Type >, 183
operator*=	operator==
BArray< Cell_Type, Data_Type >, 41 BArrayCell< Cell_Type, Data_Type >, 48 BArrayDense< Cell_Type, Data_Type >, 60 BArrayDenseCell< Cell_Type, Data_Type >, 68 BArrayRow< Cell_Type, Data_Type >, 81 BArrayVector< Cell_Type, Data_Type >, 87	BArray< Cell_Type, Data_Type >, 43 BArrayCell< Cell_Type, Data_Type >, 49 BArrayCell_const< Cell_Type, Data_Type >, 51 BArrayDense< Cell_Type, Data_Type >, 62 BArrayDenseCell< Cell_Type, Data_Type >, 69 BArrayRow< Cell_Type, Data_Type >, 82
operator()	BArrayRow_const< Cell_Type, Data_Type >, 84
BArray< Cell_Type, Data_Type >, 41 barray-meat-operators.hpp, 205 BArrayDense< Cell_Type, Data_Type >, 60 BArrayDenseCol< Cell_Type, Data_Type >, 72 BArrayDenseCol_const< Cell_Type, Data_Type >,	BArrayVector< Cell_Type, Data_Type >, 88 BArrayVector_const< Cell_Type, Data_Type >, 91 Cell< Cell_Type >, 96 operator[] Counters< Array_Type, Data_Type >, 107
74	PhyloCounterData, 167
BArrayDenseRow< Cell_Type, Data_Type >, 77 BArrayDenseRow const< Cell Type, Data Type	PowerSet< Array_Type, Data_Rule_Type >, 174 ord
>, 79	Node, 163
Flock, 115	out_of_range
PhyloCounterData, 167	BArray< Cell_Type, Data_Type >, 43
Rule < Array_Type, Data_Type >, 180	BArrayDense< Cell_Type, Data_Type >, 62
Rules< Array_Type, Data_Type >, 183	
vecHasher $<$ T $>$, 198	parent
operator+=	Node, 163
BArray< Cell_Type, Data_Type >, 41, 42	parse_polytomies
BArrayCell< Cell_Type, Data_Type >, 48	Flock, 115
BArrayDense< Cell_Type, Data_Type >, 61	Geese, 134
BArrayDenseCell< Cell_Type, Data_Type >, 69	Phylo counters, 19
BArrayRow< Cell_Type, Data_Type >, 81	counter_co_opt, 21 counter_cogain, 21
BArrayVector< Cell_Type, Data_Type >, 87	counter_gains, 21
operator-=	counter_gains_from_0, 21
BArray < Cell_Type, Data_Type >, 42	counter_gains_k_offspring, 22
BArrayCell< Cell_Type, Data_Type >, 49	counter_genes_changing, 22
BArrayDense< Cell_Type, Data_Type >, 61	counter k genes changing, 22
BArrayDenseCell< Cell_Type, Data_Type >, 69	counter_less_than_p_prop_genes_changing, 22
BArrayRow< Cell_Type, Data_Type >, 81	counter_longest, 23
BArrayVector< Cell_Type, Data_Type >, 87	counter loss, 23
operator/=	counter_maxfuns, 23
BArray< Cell_Type, Data_Type >, 42	counter_neofun, 23
BArrayCell< Cell_Type, Data_Type >, 49	counter_neofun_a2b, 24
BArrayDense < Cell_Type, Data_Type >, 62	counter_overall_changes, 24
BArrayDenseCell < Cell_Type, Data_Type >, 69	counter overall gains, 24
BArrayRow< Cell_Type, Data_Type >, 82	counter_overall_gains_from_0, 24
BArrayVector < Cell_Type, Data_Type >, 87	counter overall loss, 25
operator=	counter_pairwise_first_gain, 25
BArray Cell_Type, Data_Type >, 43	counter_pairwise_neofun_singlefun, 25
BArrayDenes < Cell Type, Data Type >, 49	counter_pairwise_overall_change, 25
BArrayDense< Cell_Type, Data_Type >, 62	counter_pairwise_preserving, 26
BArrayPow< Coll Type, Data_Type >, 69	counter_preserve_pseudogene, 26
BArrayRow< Cell_Type, Data_Type >, 82 BArrayVector< Cell_Type, Data_Type >, 88	counter_prop_genes_changing, 26
	counter_subfun, 26
Cell< Cell_Type >, 95, 96 Counter< Array_Type, Data_Type >, 102	Phylo rules, 27
Counters Array_Type, Data_Type >, 102 Counters Array_Type, Data_Type >, 106, 107	rule_dyn_limit_changes, 27
Journals Array_type, Data_type >, 100, 107	phylo.hpp

DEEN II T. BUBLIO ATION	
DEFAULT_DUPLICATION, 299	phylo.hpp, 303
DUPL_DUPL, 299	PhyloRuleDyn
DUPL_EITH, 299	phylo.hpp, 303
DUPL_SPEC, 299	PhyloRuleDynData, 168
get_last_name, 304	\sim PhyloRuleDynData, 169
IF_MATCHES, 300	counts, 169
IF_NOTMATCHES, 300	duplication, 169
IS_DUPLICATION, 300	lb, 169
IS_EITHER, 300	PhyloRuleDynData, 169
IS_SPECIATION, 300	pos, 169
MAKE_DUPL_VARS, 301	ub, 170
PHYLO_CHECK_MISSING, 301	PhyloRules
PHYLO_COUNTER_LAMBDA, 301	phylo.hpp, 303
PHYLO_RULE_DYN_LAMBDA, 301	PhyloRulesDyn
PhyloArray, 302	phylo.hpp, 303
PhyloCounter, 302	PhyloStatsCounter
PhyloCounters, 302	phylo.hpp, 303
PhyloModel, 302	PhyloSupport
PhyloPowerSet, 302	phylo.hpp, 304
PhyloRule, 303	POS
PhyloRuleData, 303	barraydense-meat-operators.hpp, 229
PhyloRuleDyn, 303	barraydense-meat.hpp, 234
PhyloRules, 303	barraydensecell-bones.hpp, 252
PhyloRulesDyn, 303	barraydensecell-meat.hpp, 254
· · · · · · · · · · · · · · · · · · ·	• • • • • • • • • • • • • • • • • • • •
PhyloStatsCounter, 303	barraydensecol-bones.hpp, 255
PhyloSupport, 304	barraydenserow-bones.hpp, 256
PHYLO_CHECK_MISSING	pos
phylo.hpp, 301	PhyloRuleDynData, 169
PHYLO_COUNTER_LAMBDA	POS_N
phylo.hpp, 301	barraydense-meat-operators.hpp, 229
PHYLO_RULE_DYN_LAMBDA	barraydense-meat.hpp, 234
phylo.hpp, 301	barraydensecol-bones.hpp, 255
PhyloArray	barraydenserow-bones.hpp, 256
phylo.hpp, 302	PowerSet
PhyloCounter	PowerSet< Array_Type, Data_Rule_Type >, 172
phylo.hpp, 302	PowerSet < Array_Type, Data_Rule_Type >, 170
PhyloCounterData, 165	\sim PowerSet, 172
at, 166	add_rule, 172, 173
begin, 166	begin, 173
empty, 166	calc, 173
end, 167	coordinates_free, 175
get_counters, 167	coordinates_locked, 175
operator(), 167	data, 175
operator[], 167	EmptyArray, 175
PhyloCounterData, 166	end, 173
push back, 167	get_data, 173
reserve, 167	get_data_ptr, 174
shrink_to_fit, 168	init_support, 174
size, 168	M, 175
PhyloCounters	N, 176
phylo.hpp, 302	n_free, 176
PhyloModel PhyloModel	n_locked, 176
phylo.hpp, 302	operator[], 174
PhyloPowerSet	PowerSet, 172
phylo.hpp, 302	
	reset, 174
PhyloRule	rules, 176
phylo.hpp, 303	rules_deleted, 176
PhyloRuleData	size, 174

predict	reset_array
Geese, 134	StatsCounter< Array_Type, Data_Type >, 187
predict_backend	Support< Array_Type, Data_Counter_Type,
Geese, 135	Data_Rule_Type, Data_Rule_Dyn_Type >,
predict_exhaust	194
Geese, 135	resize
predict_exhaust_backend	BArray< Cell_Type, Data_Type >, 44
Geese, 135, 136	barray-meat.hpp, 217
predict_sim	BArrayDense < Cell_Type, Data_Type >, 63
Geese, 136	barraydense-meat.hpp, 243, 244
print	Entries < Cell_Type >, 109
BArray< Cell_Type, Data_Type >, 43	statscounter-meat.hpp, 328
BArrayDense < Cell_Type, Data_Type >, 62	return
Flock, 116	barray-meat.hpp, 217, 222
FreqTable $<$ T $>$, 120	barraydense-meat.hpp, 250
Geese, 136	counters-meat.hpp, 281
Model< Array_Type, Data_Counter_Type,	statscounter-meat.hpp, 331
Data_Rule_Type, Data_Rule_Dyn_Type >,	support-meat.hpp, 344
151	rhs
Support< Array_Type, Data_Counter_Type,	barray-meat-operators.hpp, 205
Data_Rule_Type, Data_Rule_Dyn_Type >,	rm cell
194	BArray< Cell_Type, Data_Type >, 44
print_observed_counts	BArrayDense < Cell_Type, Data_Type >, 63
Geese, 137	barraydense-meat.hpp, 244, 245
print_stats	support-meat.hpp, 338
Model< Array_Type, Data_Counter_Type,	ROW
Data_Rule_Type, Data_Rule_Dyn_Type >,	barray-meat-operators.hpp, 203
151	barray-meat.hpp, 209, 217
printf_barry	barraydense-meat-operators.hpp, 229
barry-configuration.hpp, 263	barraydense-meat.hpp, 234
probability	row
Node, 163	BArray< Cell_Type, Data_Type >, 44
Progress, 177	BArrayDense< Cell_Type, Data_Type >, 63
~Progress, 177	row0
end, 178	barray-meat.hpp, 222
next, 178	Row_type
Progress, 177	typedefs.hpp, 348
progress.hpp	rowsum
BARRY_PROGRESS_BAR_WIDTH, 322	BArrayDense< Cell_Type, Data_Type >, 63
pset loc	Rule
Geese, 139	Rule < Array_Type, Data_Type >, 179
push_back	Rule< Array Type, Data Type >, 178
PhyloCounterData, 167	∼Rule, 179
•	D, 180
README.md, 349	operator(), 180
reduced_sequence	Rule, 179
Geese, 140	rule_dyn_limit_changes
rengine	Phylo rules, 27
Flock, 117	rule_fun_default
report	rules-bones.hpp, 323
barray-meat.hpp, 222	Rule_fun_type
barraydense-meat.hpp, 250	typedefs.hpp, 348
reserve	RULE FUNCTION
BArray< Cell_Type, Data_Type >, 43	barry.hpp, 268
BArrayDense < Cell_Type, Data_Type >, 63	defm-bones.hpp, 310
FreqTable < T >, 120	geese-bones.hpp, 313
PhyloCounterData, 167	RULE_LAMBDA
reset	barry.hpp, 268
PowerSet < Array_Type, Data_Rule_Type >, 174	2011 J. 11 pp, 200

Rules	Model< Array_Type, Data_Counter_Type,
Rules< Array_Type, Data_Type >, 181	Data_Rule_Type, Data_Rule_Dyn_Type >,
rules PowerSet< Array_Type, Data_Rule_Type >, 176	152 Support< Array_Type, Data_Counter_Type,
support-meat.hpp, 344	Data Rule Type, Data Rule Dyn Type >,
Rules< Array_Type, Data_Type >, 180	195
∼Rules, 181	set_seed
add_rule, 182	Flock, 116
get_seq, 182	Geese, 137
operator(), 183	Model < Array_Type, Data_Counter_Type,
operator=, 183	Data_Rule_Type, Data_Rule_Dyn_Type >,
Rules, 181	153
size, 183	set_transform_model
rules-bones.hpp	Model < Array_Type, Data_Counter_Type,
rule_fun_default, 323 rules_	Data_Rule_Type, Data_Rule_Dyn_Type >, 153
support-meat.hpp, 344	shrink_to_fit
rules_deleted	PhyloCounterData, 168
PowerSet< Array_Type, Data_Rule_Type >, 176	simulate
rules_dyn	Geese, 137
support-meat.hpp, 344	size
rules_zerodiag	BArrayDenseCol < Cell_Type, Data_Type >, 72
network.hpp, 296	BArrayDenseCol_const< Cell_Type, Data_Type >, 75
sample	BArrayDenseRow< Cell_Type, Data_Type >, 77
Model< Array_Type, Data_Counter_Type, Data_Rule_Type, Data_Rule_Dyn_Type >,	BArrayDenseRow_const< Cell_Type, Data_Type
151	>, 79
search	BArrayVector < Cell_Type, Data_Type >, 88
barray-meat.hpp, 223	BArrayVector_const< Cell_Type, Data_Type >, 92
sequence	Counters< Array_Type, Data_Type >, 107 FreqTable< T >, 120
Geese, 140	Model Array_Type, Data_Counter_Type,
set_counters	Data_Rule_Type, Data_Rule_Dyn_Type >,
Model < Array_Type, Data_Counter_Type,	153
Data_Rule_Type, Data_Rule_Dyn_Type >,	PhyloCounterData, 168
152	PowerSet < Array_Type, Data_Rule_Type >, 174
StatsCounter< Array_Type, Data_Type >, 187	Rules < Array_Type, Data_Type >, 183
Support < Array_Type, Data_Counter_Type,	StatsCounter< Array_Type, Data_Type >, 187
Data_Rule_Type, Data_Rule_Dyn_Type >, 194	size_unique
set_data	Model < Array_Type, Data_Counter_Type,
BArray< Cell_Type, Data_Type >, 44	Data_Rule_Type, Data_Rule_Dyn_Type >, 153
BArrayDense < Cell Type, Data Type >, 64	source
set_keygen	barray-meat.hpp, 223
Model < Array_Type, Data_Counter_Type,	barraydense-meat.hpp, 250
Data_Rule_Type, Data_Rule_Dyn_Type >,	Entries < Cell_Type >, 109
152	states
set_rengine	NodeData, 165
Model < Array_Type, Data_Counter_Type,	Statistical Models, 11
Data_Rule_Type, Data_Rule_Dyn_Type >, 152	stats_bank
set_rules	support-meat.hpp, 344
Model< Array_Type, Data_Counter_Type,	StatsCounter
Data_Rule_Type, Data_Rule_Dyn_Type >,	StatsCounter< Array_Type, Data_Type >, 184,
152	185 StatsCounter< Array_Type, Data_Type >, 184
Support< Array_Type, Data_Counter_Type,	~StatsCounter, 185
Data_Rule_Type, Data_Rule_Dyn_Type >,	add_counter, 185
194	count_all, 186
set_rules_dyn	

count current, 186	get counters, 192
count_init, 186	get counts, 192
get_counters, 186	get_current_stats, 193
get_descriptions, 186	get_data, 193
get_names, 186	get_rules, 193
reset_array, 187	get_rules_dyn, 193
set_counters, 187	hashes, 197
size, 187	hashes initialized, 197
StatsCounter, 184, 185	init_support, 193
statscounter-meat.hpp	M, 197
clear, 328	max_num_elements, 197
counter, 330	N, 197
counter_deleted, 330	n_counters, 198
counters, 330	print, 194
counters_, 330	reset_array, 194
current_stats, 330	set_counters, 194
EmptyArray, 331	set_rules, 194
f_, 331	set_rules_dyn, 195
for, 328	Support, 190
j, 331	support-meat.hpp
resize, 328	array_bank, 341
return, 331	BARRY_SUPPORT_MEAT_HPP, 335
STATSCOUNTER TEMPLATE, 327-329	calc_backend_dense, 336
STATSCOUNTER TEMPLATE ARGS, 327	calc_backend_sparse, 337
STATSCOUNTER_TYPE, 327	change_stats_different, 342
STATSCOUNTER_TEMPLATE	coord_i, 342
statscounter-meat.hpp, 327–329	coord_i, 342
	-
STATSCOUNTER_TEMPLATE_ARGS	counters, 342
statscounter-meat.hpp, 327	counters_, 342
STATSCOUNTER_TYPE	delete_counters, 342
statscounter-meat.hpp, 327	delete_rules, 343
store_psets	delete_rules_dyn, 343
Model< Array_Type, Data_Counter_Type,	else, 343
Data_Rule_Type, Data_Rule_Dyn_Type >,	f_, 343
154	for, 337
subtree_prob	hashes, 343
Node, 163	if, 337
Support	insert_cell, 338
Support< Array_Type, Data_Counter_Type,	return, 344
Data_Rule_Type, Data_Rule_Dyn_Type >,	rm_cell, 338
190	rules, 344
Support < Array_Type, Data_Counter_Type, Data_Rule_T	
Data Rule Dyn Type >, 187	rules_dyn, 344
~Support, 190	stats_bank, 344
add counter, 191	SUPPORT_TEMPLATE, 336, 338–341
add_rule, 191	SUPPORT TEMPLATE ARGS, 336
add_rule_dyn, 191	SUPPORT TYPE, 336
·	- · · · · ·
calc, 192	tmp_chng, 345
change_stats, 195	support_size
coordiantes_n_free, 195	Flock, 116
coordiantes_n_locked, 195	Geese, 138
coordinates_free, 195	Model Array_Type, Data_Counter_Type,
coordinates_locked, 196	Data_Rule_Type, Data_Rule_Dyn_Type >,
current_stats, 196	154
delete_counters, 196	SUPPORT_TEMPLATE
delete_rules, 196	support-meat.hpp, 336, 338-341
delete_rules_dyn, 196	SUPPORT_TEMPLATE_ARGS
eval_rules_dyn, 192	support-meat.hpp, 336

SUPPORT_TYPE	V
support-meat.hpp, 336	barray-meat.hpp, 223
swap_cells	barraydense-meat.hpp, 251
BArray< Cell_Type, Data_Type >, 45	va_end
BArrayDense < Cell Type, Data Type >, 64	barraydense-meat.hpp, 245
swap_cols	va_start
BArray< Cell_Type, Data_Type >, 45	barraydense-meat.hpp, 245
BArrayDense< Cell_Type, Data_Type >, 64	val
swap_rows	Entries < Cell_Type >, 110
BArray< Cell_Type, Data_Type >, 45	val0
BArrayDense < Cell_Type, Data_Type >, 64	barraydense-meat.hpp, 251
towest.	val1
target	barraydense-meat.hpp, 251
barray-meat.hpp, 223	value
barraydense-meat.hpp, 250	barray-meat.hpp, 223
Entries < Cell_Type >, 110	barraydense-meat.hpp, 251
this	Cell< Cell_Type >, 96
barray-meat-operators.hpp, 205	vec_diff
counters-meat.hpp, 281	defm-bones.hpp, 310
tmp_chng	geese-bones.hpp, 313
support-meat.hpp, 345	vec_equal
toggle_cell	typedefs.hpp, 348
BArray< Cell_Type, Data_Type >, 45	vec_equal_approx
BArrayDense< Cell_Type, Data_Type >, 65	
toggle_lock	typedefs.hpp, 349
	vec_inner_prod
BArray Cell_Type, Data_Type >, 45	typedefs.hpp, 349
BArrayDense < Cell_Type, Data_Type >, 65	vecHasher< T >, 198
transform_model	operator(), 198
Model< Array_Type, Data_Counter_Type,	vector_caster
Data_Rule_Type, Data_Rule_Dyn_Type >,	geese-bones.hpp, 314
154	vertex_attr
transpose	NetworkData, 158
BArray< Cell_Type, Data_Type >, 46	visited
BArrayDense < Cell_Type, Data_Type >, 65	BArray< Cell_Type, Data_Type >, 47
TWO	BArrayDense< Cell_Type, Data_Type >, 67
CHECK, 30	Cell< Cell_Type >, 96
EXISTS, 32	Node, 163
typedefs.hpp	vprintf
Col_type, 347	barraydense-meat.hpp, 245
Counter_fun_type, 347	barrayderise-meat.ripp, 243
Counts_type, 347	ZERO CELL
MapVec type, 348	barraydense-meat.hpp, 235
Row_type, 348	barraydensecol-bones.hpp, 255
— · ·	•
Rule_fun_type, 348	barraydenserow-bones.hpp, 256
uint, 348	zero_col
vec_equal, 348	BArray< Cell_Type, Data_Type >, 46
vec_equal_approx, 349	BArrayDense< Cell_Type, Data_Type >, 65
vec_inner_prod, 349	zero_row
	BArray< Cell_Type, Data_Type >, 46
ub	BArrayDense < Cell_Type, Data_Type >, 65
PhyloRuleDynData, 170	
uint	
typedefs.hpp, 348	
UKNOWN	
EXISTS, 32	
update_annotations	
Geese, 138	
update_normalizing_constant	
model-meat hnn 308	