Preliminary migrate analysis of M. californianus

MIGRATION RATE AND POPULATION SIZE ESTIMATION
using the coalescent and maximum likelihood or Bayesian inference
Migrate-n version 3.7.2 [April-12-18]
Program started at Tue Jun 1 13:45:41 2021
Program finished at Tue Jun 1 18:56:55 2021

Options

Datatype: DNA sequence data

Inheritance scalers in use for Thetas:
All loci use an inheritance scaler of 1.0
[The locus with a scaler of 1.0 used as reference]
Random number seed: (with internal timer) 2528787101
Start parameters:

Theta values were generated from guessed values
Theta = 0.01000
M values were generated from guessed values
M-matrix:
100000.00 [all are the same]
Connection type matrix:
where m = average (average over a group of Thetas or M,
s = symmetric M, S = symmetric 4Nm, 0 = zero, and not estimated,
* = free to vary, Thetas are on diagonal

Population	1	2	3	4	5	6	7	8	9	10	11	12
1 ElfinCo	m	m	m	m	m	m	m	m	m	m	m	m
2 Bamfiel	m	m	m	m	m	m	m	m	m	m	m	m
3 PortRen	m	m	m	m	m	m	m	m	m	m	m	m
4 WalkOnB	m	m	m	m	m	m	m	m	m	m	m	m
5 BodegaH	m	m	m	m	m	m	m	m	m	m	m	m
6 Davenpo	m	m	m	m	m	m	m	m	m	m	m	m
7 VistaDe	m	m	m	m	m	m	m	m	m	m	m	m
8 HazardR	m	m	m	m	m	m	m	m	m	m	m	m
9 Refugio	m	m	m	m	m	m	m	m	m	m	m	m
10 Carpint	m	m	m	m	m	m	m	m	m	m	m	m

11 WhitePo	*	*	*	*	*	*	*	*	*	*	*	*
12 LaJolla	*	*	*	*	*	*	*	*	*	*	*	*

Order of parameters:

1	Θ_1	=	Θ_1	[m]	<displayed>
2	Θ_2	=	Θ_1	[m]	
3	Θ_3	=	Θ_1	[m]	
4	Θ_4	=	Θ_1	[m]	
5	Θ_5	=	Θ_1	[m]	
6	Θ_6	=	Θ_1	[m]	
7	Θ_7	=	Θ_1	[m]	
8	Θ_8	=	Θ_1	[m]	
9	Θ_9	=	Θ_1	[m]	
10	Θ_{10}	=	Θ_1	[m]	
11	Θ_{11}				<displayed>
12	Θ_{12}				<displayed>
13	$M_{2\text{-->}1}$	=	$M_{2\text{-->}1}$	[m]	<displayed>
14	$M_{3\text{-->}1}$	=	$M_{2\text{-->}1}$	[m]	
15	$M_{4\text{-->}1}$	=	$M_{2\text{-->}1}$	[m]	
16	$M_{5\text{-->}1}$	=	$M_{2\text{-->}1}$	[m]	
17	$M_{6\text{-->}1}$	=	$M_{2\text{-->}1}$	[m]	
18	$M_{7\text{-->}1}$	=	$M_{2\text{-->}1}$	[m]	
19	$M_{8\text{-->}1}$	=	$M_{2\text{-->}1}$	[m]	
20	$M_{9\text{-->}1}$	=	$M_{2\text{-->}1}$	[m]	
21	$M_{10\text{-->}1}$	=	$M_{2\text{-->}1}$	[m]	
22	$M_{11\text{-->}1}$	=	$M_{2\text{-->}1}$	[m]	
23	$M_{12\text{-->}1}$	=	$M_{2\text{-->}1}$	[m]	
24	$M_{1\text{-->}2}$	=	$M_{2\text{-->}1}$	[m]	
25	$M_{3\text{-->}2}$	=	$M_{2\text{-->}1}$	[m]	
26	$M_{4\text{-->}2}$	=	$M_{2\text{-->}1}$	[m]	
27	$M_{5\text{-->}2}$	=	$M_{2\text{-->}1}$	[m]	
28	$M_{6\text{-->}2}$	=	$M_{2\text{-->}1}$	[m]	
29	$M_{7\text{-->}2}$	=	$M_{2\text{-->}1}$	[m]	
30	$M_{8\text{-->}2}$	=	$M_{2\text{-->}1}$	[m]	
31	$M_{9\text{-->}2}$	=	$M_{2\text{-->}1}$	[m]	
32	$M_{10\text{-->}2}$	=	$M_{2\text{-->}1}$	[m]	
33	$M_{11\text{-->}2}$	=	$M_{2\text{-->}1}$	[m]	
34	$M_{12\text{-->}2}$	=	$M_{2\text{-->}1}$	[m]	
35	$M_{1\text{-->}3}$	=	$M_{2\text{-->}1}$	[m]	
36	$M_{2\text{-->}3}$	=	$M_{2\text{-->}1}$	[m]	
37	$M_{4\text{-->}3}$	=	$M_{2\text{-->}1}$	[m]	
38	$M_{5\text{-->}3}$	=	$M_{2\text{-->}1}$	[m]	
39	$M_{6\text{-->}3}$	=	$M_{2\text{-->}1}$	[m]	
40	$M_{7\text{-->}3}$	=	$M_{2\text{-->}1}$	[m]	

41	$M_{8\to3}$	=	$M_{2\to1}$	[m]
42	$M_{9\to3}$	=	$M_{2\to1}$	[m]
43	$M_{10\to3}$	=	$M_{2\to1}$	[m]
44	$M_{11\to3}$	=	$M_{2\to1}$	[m]
45	$M_{12\to3}$	=	$M_{2\to1}$	[m]
46	$M_{1\to4}$	=	$M_{2\to1}$	[m]
47	$M_{2\to4}$	=	$M_{2\to1}$	[m]
48	$M_{3\to4}$	=	$M_{2\to1}$	[m]
49	$M_{5\to4}$	=	$M_{2\to1}$	[m]
50	$M_{6\to4}$	=	$M_{2\to1}$	[m]
51	$M_{7\to4}$	=	$M_{2\to1}$	[m]
52	$M_{8\to4}$	=	$M_{2\to1}$	[m]
53	$M_{9\to4}$	=	$M_{2\to1}$	[m]
54	$M_{10\to4}$	=	$M_{2\to1}$	[m]
55	$M_{11\to4}$	=	$M_{2\to1}$	[m]
56	$M_{12\to4}$	=	$M_{2\to1}$	[m]
57	$M_{1\to5}$	=	$M_{2\to1}$	[m]
58	$M_{2\to5}$	=	$M_{2\to1}$	[m]
59	$M_{3\to5}$	=	$M_{2\to1}$	[m]
60	$M_{4\to5}$	=	$M_{2\to1}$	[m]
61	$M_{6\to5}$	=	$M_{2\to1}$	[m]
62	$M_{7\to5}$	=	$M_{2\to1}$	[m]
63	$M_{8\to5}$	=	$M_{2\to1}$	[m]
64	$M_{9\to5}$	=	$M_{2\to1}$	[m]
65	$M_{10\to5}$	=	$M_{2\to1}$	[m]
66	$M_{11\to5}$	=	$M_{2\to1}$	[m]
67	$M_{12\to5}$	=	$M_{2\to1}$	[m]
68	$M_{1\to6}$	=	$M_{2\to1}$	[m]
69	$M_{2\to6}$	=	$M_{2\to1}$	[m]
70	$M_{3\to6}$	=	$M_{2\to1}$	[m]
71	$M_{4\to6}$	=	$M_{2\to1}$	[m]
72	$M_{5\to6}$	=	$M_{2\to1}$	[m]
73	$M_{7\to6}$	=	$M_{2\to1}$	[m]
74	$M_{8\to6}$	=	$M_{2\to1}$	[m]
75	$M_{9\to6}$	=	$M_{2\to1}$	[m]
76	$M_{10\to6}$	=	$M_{2\to1}$	[m]
77	$M_{11\to6}$	=	$M_{2\to1}$	[m]
78	$M_{12\to6}$	=	$M_{2\to1}$	[m]
79	$M_{1\to7}$	=	$M_{2\to1}$	[m]
80	$M_{2\to7}$	=	$M_{2\to1}$	[m]
81	$M_{3\to7}$	=	$M_{2\to1}$	[m]
82	$M_{4\to7}$	=	$M_{2\to1}$	[m]
83	$M_{5\to7}$	=	$M_{2\to1}$	[m]
84	$M_{6\to7}$	=	$M_{2\to1}$	[m]
85	$M_{8\to7}$	=	$M_{2\to1}$	[m]

86	$M_{9\rightarrow 7}$ =	$M_{2\rightarrow 1}$	[m]	
87	$M_{10\rightarrow 7}$ =	$M_{2\rightarrow 1}$	[m]	
88	$M_{11\rightarrow 7}$ =	$M_{2\rightarrow 1}$	[m]	
89	$M_{12\rightarrow 7}$ =	$M_{2\rightarrow 1}$	[m]	
90	$M_{1\rightarrow 8}$ =	$M_{2\rightarrow 1}$	[m]	
91	$M_{2\rightarrow 8}$ =	$M_{2\rightarrow 1}$	[m]	
92	$M_{3\rightarrow 8}$ =	$M_{2\rightarrow 1}$	[m]	
93	$M_{4\rightarrow 8}$ =	$M_{2\rightarrow 1}$	[m]	
94	$M_{5\rightarrow 8}$ =	$M_{2\rightarrow 1}$	[m]	
95	$M_{6\rightarrow 8}$ =	$M_{2\rightarrow 1}$	[m]	
96	$M_{7\rightarrow 8}$ =	$M_{2\rightarrow 1}$	[m]	
97	$M_{9\rightarrow 8}$ =	$M_{2\rightarrow 1}$	[m]	
98	$M_{10\rightarrow 8}$ =	$M_{2\rightarrow 1}$	[m]	
99	$M_{11\rightarrow 8}$ =	$M_{2\rightarrow 1}$	[m]	
100	$M_{12\rightarrow 8}$ =	$M_{2\rightarrow 1}$	[m]	
101	$M_{1\rightarrow 9}$ =	$M_{2\rightarrow 1}$	[m]	
102	$M_{2\rightarrow 9}$ =	$M_{2\rightarrow 1}$	[m]	
103	$M_{3\rightarrow 9}$ =	$M_{2\rightarrow 1}$	[m]	
104	$M_{4\rightarrow 9}$ =	$M_{2\rightarrow 1}$	[m]	
105	$M_{5\rightarrow 9}$ =	$M_{2\rightarrow 1}$	[m]	
106	$M_{6\rightarrow 9}$ =	$M_{2\rightarrow 1}$	[m]	
107	$M_{7\rightarrow 9}$ =	$M_{2\rightarrow 1}$	[m]	
108	$M_{8\rightarrow 9}$ =	$M_{2\rightarrow 1}$	[m]	
109	$M_{10\rightarrow 9}$ =	$M_{2\rightarrow 1}$	[m]	
110	$M_{11\rightarrow 9}$ =	$M_{2\rightarrow 1}$	[m]	
111	$M_{12\rightarrow 9}$ =	$M_{2\rightarrow 1}$	[m]	
112	$M_{1\rightarrow 10}$ =	$M_{2\rightarrow 1}$	[m]	
113	$M_{2\rightarrow 10}$ =	$M_{2\rightarrow 1}$	[m]	
114	$M_{3\rightarrow 10}$ =	$M_{2\rightarrow 1}$	[m]	
115	$M_{4\rightarrow 10}$ =	$M_{2\rightarrow 1}$	[m]	
116	$M_{5\rightarrow 10}$ =	$M_{2\rightarrow 1}$	[m]	
117	$M_{6\rightarrow 10}$ =	$M_{2\rightarrow 1}$	[m]	
118	$M_{7\rightarrow 10}$ =	$M_{2\rightarrow 1}$	[m]	
119	$M_{8\rightarrow 10}$ =	$M_{2\rightarrow 1}$	[m]	
120	$M_{9\rightarrow 10}$ =	$M_{2\rightarrow 1}$	[m]	
121	$M_{11\rightarrow 10}$ =	$M_{2\rightarrow 1}$	[m]	
122	$M_{12\rightarrow 10}$ =	$M_{2\rightarrow 1}$	[m]	
123	$M_{1\rightarrow 11}$			<displayed>
124	$M_{2\rightarrow 11}$			<displayed>
125	$M_{3\rightarrow 11}$			<displayed>
126	$M_{4\rightarrow 11}$			<displayed>
127	$M_{5\rightarrow 11}$			<displayed>
128	$M_{6\rightarrow 11}$			<displayed>
129	$M_{7\rightarrow 11}$			<displayed>
130	$M_{8\rightarrow 11}$			<displayed>

131	$M_{9->11}$	<displayed>
132	$M_{10->11}$	<displayed>
133	$M_{12->11}$	<displayed>
134	$M_{1->12}$	<displayed>
135	$M_{2->12}$	<displayed>
136	$M_{3->12}$	<displayed>
137	$M_{4->12}$	<displayed>
138	$M_{5->12}$	<displayed>
139	$M_{6->12}$	<displayed>
140	$M_{7->12}$	<displayed>
141	$M_{8->12}$	<displayed>
142	$M_{9->12}$	<displayed>
143	$M_{10->12}$	<displayed>
144	$M_{11->12}$	<displayed>

Mutation rate among loci: Mutation rate is constant

Analysis strategy: Bayesian inference

Proposal distributions for parameter

Parameter	Proposal
Theta	Metropolis sampling
M	Slice sampling

Prior distribution for parameter

Parameter	Prior	Minimum	Mean*	Maximum	Delta	Bins
Theta	Exp window	0.000010	0.010000	10.000000	1.000000	500
M	Exp window	0.000100	100000.000000	1000000.000000	100000.000000	500

Markov chain settings:	Long chain
Number of chains	1
Recorded steps [a]	1000
Increment (record every x step [b]	100
Number of concurrent chains (replicates) [c]	3
Visited (sampled) parameter values [a*b*c]	300000
Number of discard trees per chain (burn-in)	1000

Multiple Markov chains:

Static heating scheme — 4 chains with temperatures

100000.00 3.00 1.50 1.00

Swapping interval is 1

Print options:

Data file:	../../mcalifornianus_210528.mig
Output file:	outfile.txt
Posterior distribution raw histogram file:	bayesfile
Print data:	No
Print genealogies [only some for some data type]:	None

Data summary

Datatype:	Sequence data
Number of loci:	1

Population	Locus	Gene copies
1 ElfinCo	1	19
2 Bamfiel	1	23
3 PortRen	1	15
4 WalkOnB	1	16
5 BodegaH	1	7
6 Davenpo	1	17
7 VistaDe	1	19
8 HazardR	1	23
9 Refugio	1	16
10 Carpint	1	19
11 WhitePo	1	11
12 LaJolla	1	8
Total of all populations	1	193

Bayesian Analysis: Posterior distribution table

Locus	Parameter	2.5%	25.0%	Mode	75.0%	97.5%	Median	Mean
1	Θ_1	0.00001	0.00001	0.01001	0.06001	0.16001	0.07001	0.00076
1	Θ_2	0.00001	0.00001	0.01001	0.06001	0.16001	0.07001	0.00076
1	Θ_3	0.00001	0.00001	0.01001	0.06001	0.16001	0.07001	0.00076
1	Θ_4	0.00001	0.00001	0.01001	0.06001	0.16001	0.07001	0.00076
1	Θ_5	0.00001	0.00001	0.01001	0.06001	0.16001	0.07001	0.00076
1	Θ_6	0.00001	0.00001	0.01001	0.06001	0.16001	0.07001	0.00076
1	Θ_7	0.00001	0.00001	0.01001	0.06001	0.16001	0.07001	0.00076
1	Θ_8	0.00001	0.00001	0.01001	0.06001	0.16001	0.07001	0.00076
1	Θ_9	0.00001	0.00001	0.01001	0.06001	0.16001	0.07001	0.00076
1	Θ_{10}	0.00001	0.00001	0.01001	0.06001	0.16001	0.07001	0.00076
1	Θ_{11}	0.00001	0.00001	0.01001	0.06001	0.16001	0.07001	0.01013
1	Θ_{12}	0.00001	0.00001	0.01001	0.06001	0.16001	0.07001	0.00976
1	$M_{2->1}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{3->1}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{4->1}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{5->1}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{6->1}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{7->1}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{8->1}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{9->1}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{10->1}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{11->1}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{12->1}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{1->2}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{3->2}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{4->2}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{5->2}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{6->2}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{7->2}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{8->2}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{9->2}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{10->2}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{11->2}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{12->2}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{1->3}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{2->3}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8

Locus	Parameter	2.5%	25.0%	Mode	75.0%	97.5%	Median	Mean
1	$M_{4->3}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{5->3}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{6->3}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{7->3}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{8->3}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{9->3}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{10->3}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{11->3}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{12->3}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{1->4}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{2->4}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{3->4}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{5->4}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{6->4}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{7->4}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{8->4}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{9->4}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{10->4}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{11->4}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{12->4}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{1->5}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{2->5}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{3->5}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{4->5}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{6->5}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{7->5}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{8->5}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{9->5}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{10->5}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{11->5}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{12->5}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{1->6}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{2->6}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{3->6}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{4->6}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{5->6}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{7->6}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{8->6}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{9->6}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{10->6}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{11->6}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8

Locus	Parameter	2.5%	25.0%	Mode	75.0%	97.5%	Median	Mean
1	$M_{12->6}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{1->7}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{2->7}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{3->7}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{4->7}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{5->7}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{6->7}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{8->7}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{9->7}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{10->7}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{11->7}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{12->7}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{1->8}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{2->8}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{3->8}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{4->8}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{5->8}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{6->8}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{7->8}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{9->8}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{10->8}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{11->8}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{12->8}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{1->9}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{2->9}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{3->9}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{4->9}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{5->9}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{6->9}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{7->9}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{8->9}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{10->9}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{11->9}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{12->9}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{1->10}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{2->10}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{3->10}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{4->10}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{5->10}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{6->10}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{7->10}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8

Locus	Parameter	2.5%	25.0%	Mode	75.0%	97.5%	Median	Mean
1	$M_{8->10}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{9->10}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{11->10}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{12->10}$	26000.0	40000.0	49000.0	58000.0	70000.0	51000.0	49514.8
1	$M_{1->11}$	120000.0	136000.0	153000.0	170000.0	318000.0	217000.0	215006.4
1	$M_{2->11}$	0.0	0.0	1000.0	46000.0	122000.0	49000.0	53264.0
1	$M_{3->11}$	32000.0	50000.0	85000.0	118000.0	134000.0	273000.0	289159.6
1	$M_{4->11}$	44000.0	190000.0	231000.0	274000.0	348000.0	225000.0	232084.3
1	$M_{5->11}$	24000.0	108000.0	127000.0	144000.0	176000.0	107000.0	101613.2
1	$M_{6->11}$	36000.0	56000.0	87000.0	128000.0	178000.0	119000.0	238019.6
1	$M_{7->11}$	68000.0	104000.0	147000.0	166000.0	194000.0	117000.0	103700.5
1	$M_{8->11}$	12000.0	28000.0	53000.0	74000.0	108000.0	69000.0	128065.1
1	$M_{9->11}$	4000.0	16000.0	69000.0	92000.0	236000.0	177000.0	213406.2
1	$M_{10->11}$	50000.0	64000.0	89000.0	120000.0	284000.0	161000.0	161627.9
1	$M_{12->11}$	44000.0	62000.0	81000.0	134000.0	174000.0	127000.0	173800.5
1	$M_{1->12}$	18000.0	32000.0	47000.0	62000.0	80000.0	139000.0	127716.9
1	$M_{2->12}$	84000.0	168000.0	203000.0	242000.0	302000.0	203000.0	199354.1
1	$M_{3->12}$	56000.0	66000.0	87000.0	108000.0	272000.0	155000.0	153496.8
1	$M_{4->12}$	142000.0	166000.0	189000.0	228000.0	306000.0	217000.0	221585.3
1	$M_{5->12}$	46000.0	68000.0	123000.0	150000.0	178000.0	139000.0	176559.1
1	$M_{6->12}$	28000.0	46000.0	63000.0	80000.0	132000.0	71000.0	73782.5
1	$M_{7->12}$	0.0	16000.0	61000.0	80000.0	100000.0	73000.0	151841.4
1	$M_{8->12}$	16000.0	86000.0	117000.0	142000.0	170000.0	137000.0	191729.4
1	$M_{9->12}$	30000.0	54000.0	71000.0	96000.0	226000.0	123000.0	123697.1
1	$M_{10->12}$	20000.0	82000.0	113000.0	148000.0	200000.0	115000.0	113307.9
1	$M_{11->12}$	2000.0	26000.0	53000.0	66000.0	104000.0	53000.0	53375.1

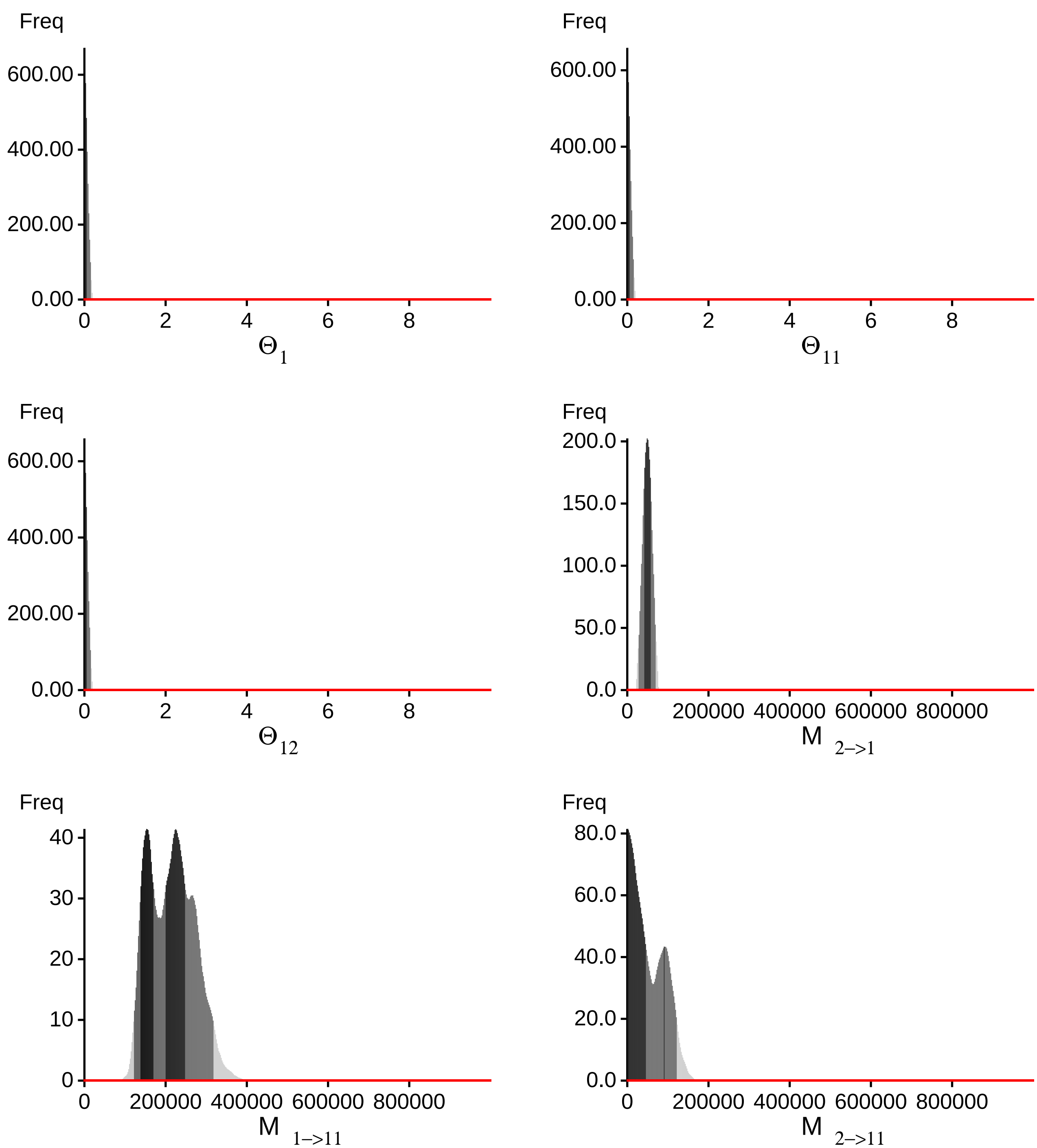
Citation suggestions:

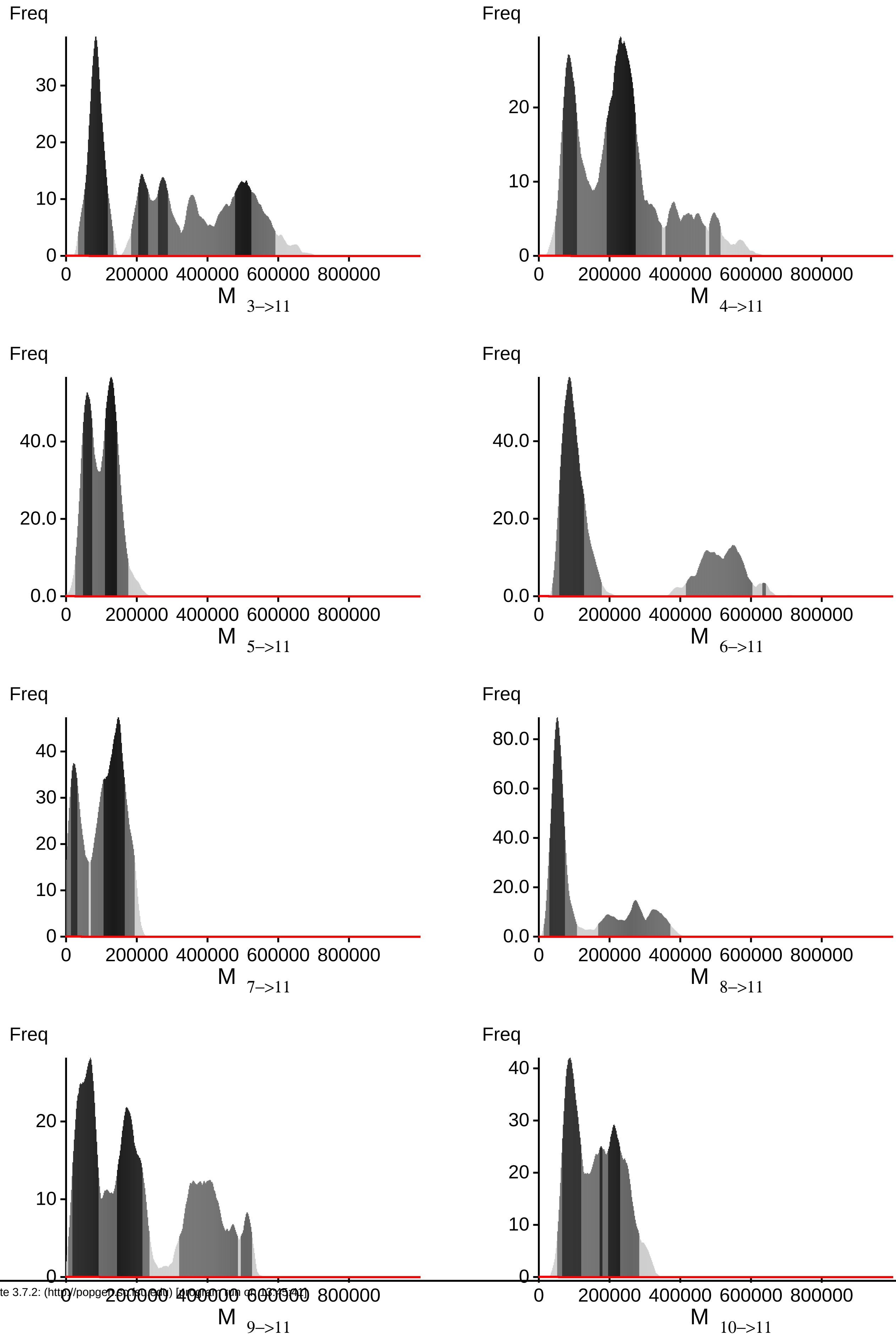
Beerli P., 2006. Comparison of Bayesian and maximum-likelihood inference of population genetic parameters. Bioinformatics 22:341-345

Beerli P., 2007. Estimation of the population scaled mutation rate from microsatellite data, Genetics, 177:1967-1968.

Beerli P., 2009. How to use MIGRATE or why are Markov chain Monte Carlo programs difficult to use? In Population Genetics for Animal Conservation, G. Bertorelle, M. W. Bruford, H. C. Hauffe, A. Rizzoli, and C. Vernesi, eds., vol. 17 of Conservation Biology, Cambridge University Press, Cambridge UK, pp. 42-79.

Bayesian Analysis: Posterior distribution over all loci

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Freq
80.0
60.0
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20.0
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M 8–>11
Freq
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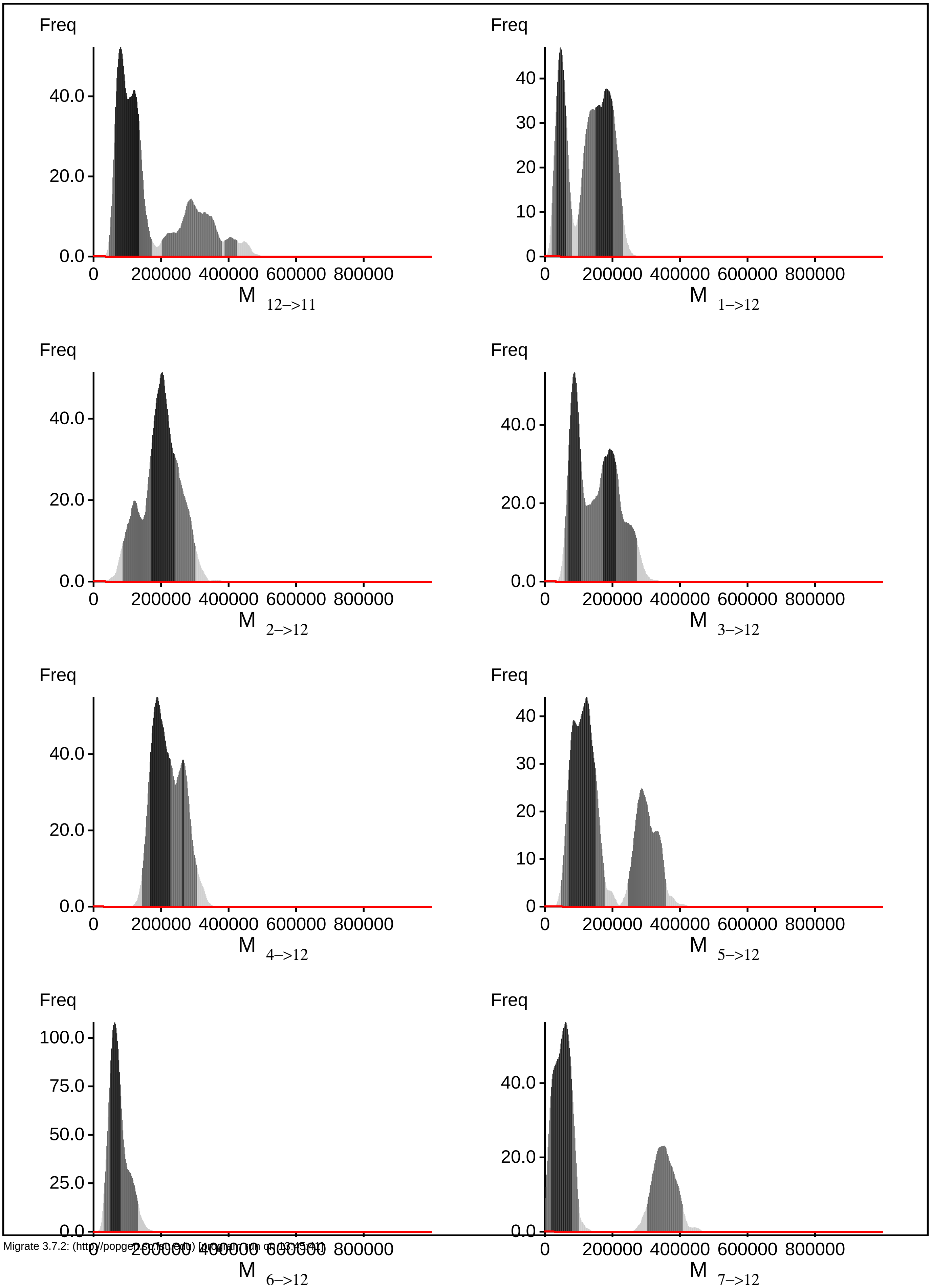
M 10–>11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M 6–>12
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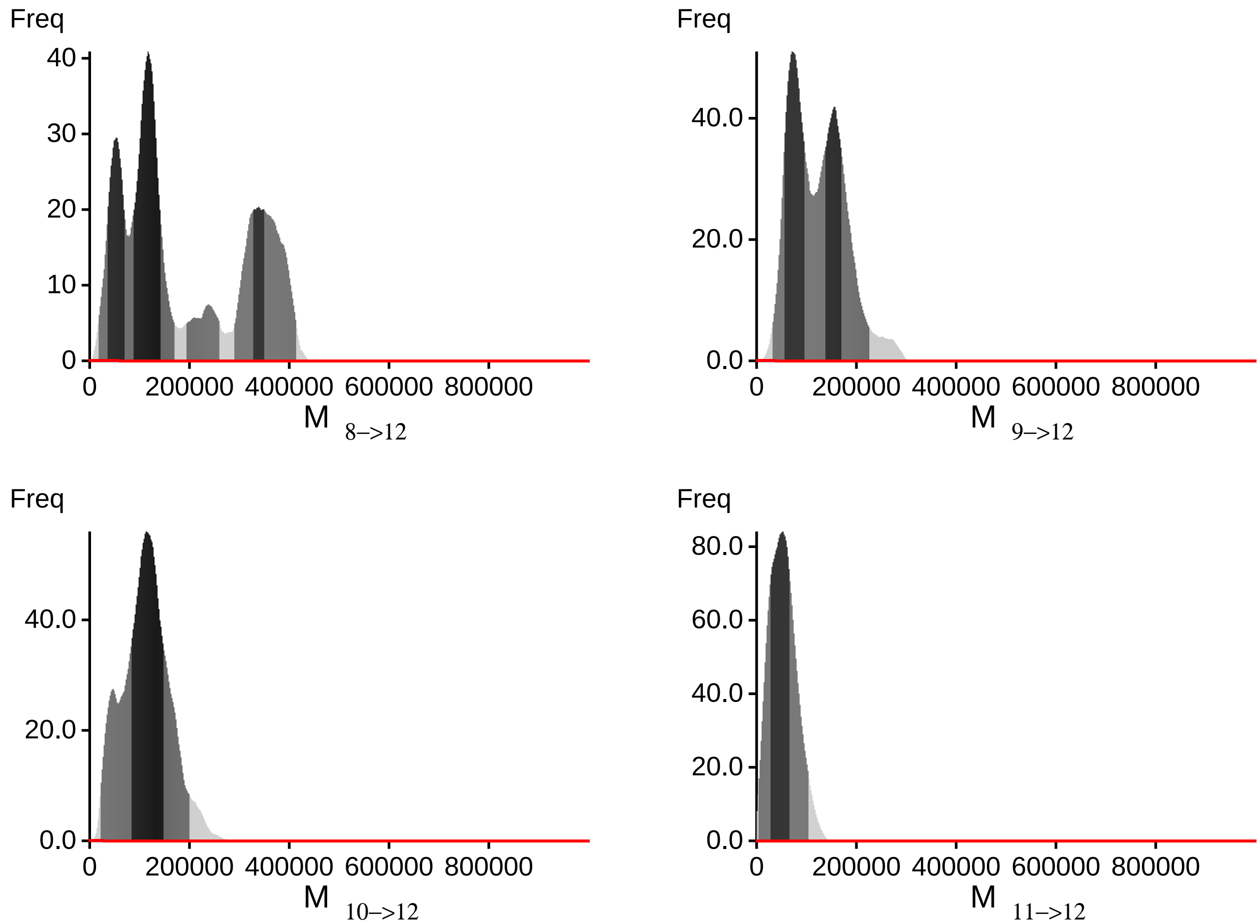
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M 9–>12
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20.0
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200000 400000 600000 800000
M 10–>12
Freq
80.0
60.0
40.0
20.0
0.0
0
200000 400000 600000 800000
M 11–>12

Log-Probability of the data given the model (marginal likelihood)

Use this value for Bayes factor calculations:
BF = Exp[ln(Prob(D | thisModel) - ln(Prob(D | otherModel)
or as LBF = 2 (ln(Prob(D | thisModel) - ln(Prob(D | otherModel))
shows the support for thisModel]

Method	ln(Prob(D|Model))	Notes
Thermodynamic integration	-2383.237361	(1a)
	-2263.537054	(1b)
Harmonic mean	-1963.481837	(2)

(1a, 1b and 2) are approximations to the marginal likelihood, make sure that the program run long enough!
(1a, 1b) and (2) should give similar results, in principle.
But (2) is overestimating the likelihood, it is presented for historical reasons and should not be used
(1a, 1b) needs heating with chains that span a temperature range of 1.0 to at least 100,000.
(1b) is using a Bezier-curve to get better approximations for runs with low number of heated chains

Citation suggestions:

Beerli P. and M. Palczewski, 2010. Unified framework to evaluate panmixia and migration direction among multiple sampling locations, Genetics, 185: 313-326.

Acceptance ratios for all parameters and the genealogies

Parameter	Accepted changes	Ratio
Θ_1	15/1038	0.01445
Θ_2	15/1038	0.01445
Θ_3	15/1038	0.01445
Θ_4	15/1038	0.01445
Θ_5	15/1038	0.01445
Θ_6	15/1038	0.01445
Θ_7	15/1038	0.01445
Θ_8	15/1038	0.01445
Θ_9	15/1038	0.01445
Θ_{10}	15/1038	0.01445
Θ_{11}	779/1036	0.75193
Θ_{12}	730/1057	0.69063
$M_{2->1}$	1049/1049	1.00000
$M_{3->1}$	1049/1049	1.00000
$M_{4->1}$	1049/1049	1.00000
$M_{5->1}$	1049/1049	1.00000
$M_{6->1}$	1049/1049	1.00000
$M_{7->1}$	1049/1049	1.00000
$M_{8->1}$	1049/1049	1.00000
$M_{9->1}$	1049/1049	1.00000
$M_{10->1}$	1049/1049	1.00000
$M_{11->1}$	1049/1049	1.00000
$M_{12->1}$	1049/1049	1.00000
$M_{1->2}$	1049/1049	1.00000
$M_{3->2}$	1049/1049	1.00000
$M_{4->2}$	1049/1049	1.00000
$M_{5->2}$	1049/1049	1.00000
$M_{6->2}$	1049/1049	1.00000
$M_{7->2}$	1049/1049	1.00000
$M_{8->2}$	1049/1049	1.00000
$M_{9->2}$	1049/1049	1.00000
$M_{10->2}$	1049/1049	1.00000
$M_{11->2}$	1049/1049	1.00000
$M_{12->2}$	1049/1049	1.00000
$M_{1->3}$	1049/1049	1.00000
$M_{2->3}$	1049/1049	1.00000
$M_{4->3}$	1049/1049	1.00000

$M_{5\to3}$	1049/1049	1.00000
$M_{6\to3}$	1049/1049	1.00000
$M_{7\to3}$	1049/1049	1.00000
$M_{8\to3}$	1049/1049	1.00000
$M_{9\to3}$	1049/1049	1.00000
$M_{10\to3}$	1049/1049	1.00000
$M_{11\to3}$	1049/1049	1.00000
$M_{12\to3}$	1049/1049	1.00000
$M_{1\to4}$	1049/1049	1.00000
$M_{2\to4}$	1049/1049	1.00000
$M_{3\to4}$	1049/1049	1.00000
$M_{5\to4}$	1049/1049	1.00000
$M_{6\to4}$	1049/1049	1.00000
$M_{7\to4}$	1049/1049	1.00000
$M_{8\to4}$	1049/1049	1.00000
$M_{9\to4}$	1049/1049	1.00000
$M_{10\to4}$	1049/1049	1.00000
$M_{11\to4}$	1049/1049	1.00000
$M_{12\to4}$	1049/1049	1.00000
$M_{1\to5}$	1049/1049	1.00000
$M_{2\to5}$	1049/1049	1.00000
$M_{3\to5}$	1049/1049	1.00000
$M_{4\to5}$	1049/1049	1.00000
$M_{6\to5}$	1049/1049	1.00000
$M_{7\to5}$	1049/1049	1.00000
$M_{8\to5}$	1049/1049	1.00000
$M_{9\to5}$	1049/1049	1.00000
$M_{10\to5}$	1049/1049	1.00000
$M_{11\to5}$	1049/1049	1.00000
$M_{12\to5}$	1049/1049	1.00000
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$M_{2\to6}$	1049/1049	1.00000
$M_{3\to6}$	1049/1049	1.00000
$M_{4\to6}$	1049/1049	1.00000
$M_{5\to6}$	1049/1049	1.00000
$M_{7\to6}$	1049/1049	1.00000
$M_{8\to6}$	1049/1049	1.00000
$M_{9\to6}$	1049/1049	1.00000
$M_{10\to6}$	1049/1049	1.00000
$M_{11\to6}$	1049/1049	1.00000
$M_{12\to6}$	1049/1049	1.00000
$M_{1\to7}$	1049/1049	1.00000
$M_{2\to7}$	1049/1049	1.00000
$M_{3\to7}$	1049/1049	1.00000
$M_{4\to7}$	1049/1049	1.00000

$M_{5->7}$	1049/1049	1.00000
$M_{6->7}$	1049/1049	1.00000
$M_{8->7}$	1049/1049	1.00000
$M_{9->7}$	1049/1049	1.00000
$M_{10->7}$	1049/1049	1.00000
$M_{11->7}$	1049/1049	1.00000
$M_{12->7}$	1049/1049	1.00000
$M_{1->8}$	1049/1049	1.00000
$M_{2->8}$	1049/1049	1.00000
$M_{3->8}$	1049/1049	1.00000
$M_{4->8}$	1049/1049	1.00000
$M_{5->8}$	1049/1049	1.00000
$M_{6->8}$	1049/1049	1.00000
$M_{7->8}$	1049/1049	1.00000
$M_{9->8}$	1049/1049	1.00000
$M_{10->8}$	1049/1049	1.00000
$M_{11->8}$	1049/1049	1.00000
$M_{12->8}$	1049/1049	1.00000
$M_{1->9}$	1049/1049	1.00000
$M_{2->9}$	1049/1049	1.00000
$M_{3->9}$	1049/1049	1.00000
$M_{4->9}$	1049/1049	1.00000
$M_{5->9}$	1049/1049	1.00000
$M_{6->9}$	1049/1049	1.00000
$M_{7->9}$	1049/1049	1.00000
$M_{8->9}$	1049/1049	1.00000
$M_{10->9}$	1049/1049	1.00000
$M_{11->9}$	1049/1049	1.00000
$M_{12->9}$	1049/1049	1.00000
$M_{1->10}$	1049/1049	1.00000
$M_{2->10}$	1049/1049	1.00000
$M_{3->10}$	1049/1049	1.00000
$M_{4->10}$	1049/1049	1.00000
$M_{5->10}$	1049/1049	1.00000
$M_{6->10}$	1049/1049	1.00000
$M_{7->10}$	1049/1049	1.00000
$M_{8->10}$	1049/1049	1.00000
$M_{9->10}$	1049/1049	1.00000
$M_{11->10}$	1049/1049	1.00000
$M_{12->10}$	1049/1049	1.00000
$M_{1->11}$	1066/1066	1.00000
$M_{2->11}$	1032/1032	1.00000
$M_{3->11}$	1041/1041	1.00000
$M_{4->11}$	1019/1019	1.00000
$M_{5->11}$	1035/1035	1.00000

$M_{6\to 11}$	1022/1022	1.00000
$M_{7\to 11}$	1047/1047	1.00000
$M_{8\to 11}$	1041/1041	1.00000
$M_{9\to 11}$	1026/1026	1.00000
$M_{10\to 11}$	1041/1041	1.00000
$M_{12\to 11}$	1044/1044	1.00000
$M_{1\to 12}$	1079/1079	1.00000
$M_{2\to 12}$	1122/1122	1.00000
$M_{3\to 12}$	1078/1078	1.00000
$M_{4\to 12}$	1056/1056	1.00000
$M_{5\to 12}$	1034/1034	1.00000
$M_{6\to 12}$	1067/1067	1.00000
$M_{7\to 12}$	1045/1045	1.00000
$M_{8\to 12}$	1014/1014	1.00000
$M_{9\to 12}$	1011/1011	1.00000
$M_{10\to 12}$	1063/1063	1.00000
$M_{11\to 12}$	995/995	1.00000
Genealogies	24140/150049	0.16088

MCMC-Autocorrelation and Effective MCMC Sample Size

Parameter	Autocorrelation	Effective Sampe Size
Θ_1	0.99076	13.91
Θ_2	0.99076	13.91
Θ_3	0.99076	13.91
Θ_4	0.99076	13.91
Θ_5	0.99076	13.91
Θ_6	0.99076	13.91
Θ_7	0.99076	13.91
Θ_8	0.99076	13.91
Θ_9	0.99076	13.91
Θ_{10}	0.99076	13.91
Θ_{11}	0.81981	298.90
Θ_{12}	0.80744	320.87
$M_{2\text{->}1}$	0.98924	16.21
$M_{3\text{->}1}$	0.98924	16.21
$M_{4\text{->}1}$	0.98924	16.21
$M_{5\text{->}1}$	0.98924	16.21
$M_{6\text{->}1}$	0.98924	16.21
$M_{7\text{->}1}$	0.98924	16.21
$M_{8\text{->}1}$	0.98924	16.21
$M_{9\text{->}1}$	0.98924	16.21
$M_{10\text{->}1}$	0.98924	16.21
$M_{11\text{->}1}$	0.98924	16.21
$M_{12\text{->}1}$	0.98924	16.21
$M_{1\text{->}2}$	0.98924	16.21
$M_{3\text{->}2}$	0.98924	16.21
$M_{4\text{->}2}$	0.98924	16.21
$M_{5\text{->}2}$	0.98924	16.21
$M_{6\text{->}2}$	0.98924	16.21
$M_{7\text{->}2}$	0.98924	16.21
$M_{8\text{->}2}$	0.98924	16.21
$M_{9\text{->}2}$	0.98924	16.21
$M_{10\text{->}2}$	0.98924	16.21
$M_{11\text{->}2}$	0.98924	16.21
$M_{12\text{->}2}$	0.98924	16.21
$M_{1\text{->}3}$	0.98924	16.21
$M_{2\text{->}3}$	0.98924	16.21
$M_{4\text{->}3}$	0.98924	16.21

$M_{5\to3}$	0.98924	16.21
$M_{6\to3}$	0.98924	16.21
$M_{7\to3}$	0.98924	16.21
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$M_{12\to6}$	0.98924	16.21
$M_{1\to7}$	0.98924	16.21
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$M_{5\to7}$	0.98924	16.21
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$M_{9\to7}$	0.98924	16.21
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$M_{11\to7}$	0.98924	16.21
$M_{12\to7}$	0.98924	16.21
$M_{1\to8}$	0.98924	16.21
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$M_{12\to8}$	0.98924	16.21
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$M_{12\to9}$	0.98924	16.21
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$M_{8\to10}$	0.98924	16.21
$M_{9\to10}$	0.98924	16.21
$M_{11\to10}$	0.98924	16.21
$M_{12\to10}$	0.98924	16.21
$M_{1\to11}$	0.87882	194.08
$M_{2\to11}$	0.91056	140.52
$M_{3\to11}$	0.93347	103.47
$M_{4\to11}$	0.94481	85.59
$M_{5\to11}$	0.88843	183.64

$M_{6\rightarrow 11}$	0.91020	142.16
$M_{7\rightarrow 11}$	0.88996	177.04
$M_{8\rightarrow 11}$	0.91120	139.56
$M_{9\rightarrow 11}$	0.93967	93.53
$M_{10\rightarrow 11}$	0.88455	185.89
$M_{12\rightarrow 11}$	0.91447	135.09
$M_{1\rightarrow 12}$	0.90042	157.49
$M_{2\rightarrow 12}$	0.90508	150.76
$M_{3\rightarrow 12}$	0.91531	133.38
$M_{4\rightarrow 12}$	0.90576	149.59
$M_{5\rightarrow 12}$	0.91496	134.29
$M_{6\rightarrow 12}$	0.84943	245.64
$M_{7\rightarrow 12}$	0.89487	167.39
$M_{8\rightarrow 12}$	0.93228	106.94
$M_{9\rightarrow 12}$	0.92583	116.99
$M_{10\rightarrow 12}$	0.94658	82.75
$M_{11\rightarrow 12}$	0.91161	138.78
Ln[Prob(D|G)]	0.98292	25.84

Potential Problems

This section reports potential problems with your run, but such reporting is often not very accurate. Whith many parameters in a multilocus analysis, it is very common that some parameters for some loci will not be very informative, triggering suggestions (for example to increase the prior range) that are not sensible. This suggestion tool will improve with time, therefore do not blindly follow its suggestions. If some parameters are flagged, inspect the tables carefully and judge wether an action is required. For example, if you run a Bayesian inference with sequence data, for macroscopic species there is rarely the need to increase the prior for Theta beyond 0.1; but if you use microsatellites it is rather common that your prior distribution for Theta should have a range from 0.0 to 100 or more. With many populations (>3) it is also very common that some migration routes are estimated poorly because the data contains little or no information for that route. Increasing the range will not help in such situations, reducing number of parameters may help in such situations.

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