## **AUTO**

POPULATION SIZE, MIGRATION, DIVERGENCE, ASSIGNMENT, HISTORY

Bayesian inference using the structured coalescent

Migrate-n version 5.0.0a [May-20-2017]

Using Intel AVX (Advanced Vector Extensions)

Compiled for PARALLEL computer architectures

One master and 100 compute nodes are available.

Program started at Sat Aug 12 19:59:31 2017

Program finished at Sat Aug 12 21:31:23 2017 [Runtime:0000:01:31:52]



### **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) 4142380767

Start parameters:

Theta values were generated Using a percent value of the prior

M values were generated Using a percent value of the prior

Connection matrix:

m = average (average over a group of Thetas or M,

s = symmetric migration M, S = symmetric 4Nm,

0 = zero, and not estimated,

\* = migration free to vary, Thetas are on diagonal

1

d = row population split off column population, D = split and then migration

Population

1 Romanshorn 0 \*

Order of parameters:

1  $\Theta_1$  <displayed>

Mutation rate among loci: Mutation rate is constant for all loci

Analysis strategy:

Bayesian inference

-Population size estimation: Exponential Distribution

Proposal distributions for parameter

Parameter Proposal
Theta Metropolis sampling
M Metropolis sampling
Divergence Metropolis sampling
Divergence Spread Metropolis sampling
Genealogy Metropolis-Hastings

Prior distribution for parameter

Parameter Prior Minimum MeanMaximum Delta Bins UpdateFreq
1 Theta -11 Uniform 0.000000 0.050 0.100 0.010 1500 0.20000

[-1 -1 means priors were set globally]

Number of concurrent chains (replicates) [c]

Markov chain settings:

Long chain

Number of chains1Recorded steps [a]50000Increment (record every x step [b]200

Visited (sampled) parameter values [a\*b\*c] 20000000

Number of discard trees per chain (burn-in) 10000

Multiple Markov chains:

Static heating scheme 4 chains with temperatures

1000000.00 3.00 1.50 1.00

Swapping interval is 1

Print options:

Data file: infile.0.9

Haplotyping is turned on:

Output file: outfile\_0.9\_0.5

Posterior distribution raw histogram file: bayesfile
Raw data from the MCMC run: bayesallfile\_0.9\_0.5

Print data: No

Print genealogies [only some for some data type]:

# Data summary

Data file: infile.0.9
Datatype: Sequence data
Number of loci: 100

N 4.				I	- 1	_
11//11	utat	ınr	١m	റവ	ρı	•

Mutation	model:			
Locus S	ublocus	Mutationmodel	Mutationmodel parameters	
1	1	Jukes-Cantor	[Basefreq: =0.25]	
2	1	Jukes-Cantor	[Basefreq: =0.25]	
3	1	Jukes-Cantor	[Basefreq: =0.25]	
4	1	Jukes-Cantor	[Basefreq: =0.25]	
5	1	Jukes-Cantor	[Basefreq: =0.25]	
6	1	Jukes-Cantor	[Basefreq: =0.25]	
7	1	Jukes-Cantor	[Basefreq: =0.25]	
8	1	Jukes-Cantor	[Basefreq: =0.25]	
9	1	Jukes-Cantor	[Basefreq: =0.25]	
10	1	Jukes-Cantor	[Basefreq: =0.25]	
11	1	Jukes-Cantor	[Basefreq: =0.25]	
12	1	Jukes-Cantor	[Basefreq: =0.25]	
13	1	Jukes-Cantor	[Basefreq: =0.25]	
14	1	Jukes-Cantor	[Basefreq: =0.25]	
15	1	Jukes-Cantor	[Basefreq: =0.25]	
16	1	Jukes-Cantor	[Basefreq: =0.25]	
17	1	Jukes-Cantor	[Basefreq: =0.25]	
18	1	Jukes-Cantor	[Basefreq: =0.25]	
19	1	Jukes-Cantor	[Basefreq: =0.25]	
20	1	Jukes-Cantor	[Basefreq: =0.25]	
21	1	Jukes-Cantor	[Basefreq: =0.25]	
22	1	Jukes-Cantor	[Basefreq: =0.25]	
23	1	Jukes-Cantor	[Basefreq: =0.25]	
24	1	Jukes-Cantor	[Basefreq: =0.25]	
25	1	Jukes-Cantor	[Basefreq: =0.25]	
26	1	Jukes-Cantor	[Basefreq: =0.25]	
27	1	Jukes-Cantor	[Basefreq: =0.25]	
28	1	Jukes-Cantor	[Basefreq: =0.25]	
29	1	Jukes-Cantor	[Basefreq: =0.25]	
30	1	Jukes-Cantor	[Basefreq: =0.25]	
31	1	Jukes-Cantor	[Basefreq: =0.25]	
32	1	Jukes-Cantor	[Basefreq: =0.25]	
33	1	Jukes-Cantor	[Basefreq: =0.25]	
34	1	Jukes-Cantor	[Basefreq: =0.25]	

25	1	lukoa Cantar	[Pagefreg: _0.25]
35 36	1 1	Jukes-Cantor Jukes-Cantor	[Basefreq: =0.25]
37	1	Jukes-Cantor	[Basefreq: =0.25] [Basefreq: =0.25]
38	1	Jukes-Cantor	[Basefreq: =0.25]
39	1	Jukes-Cantor	[Basefreq: =0.25]
40	1	Jukes-Cantor	[Basefreq: =0.25]
41	1	Jukes-Cantor	[Basefreq: =0.25]
42	1	Jukes-Cantor	[Basefreq: =0.25]
43	1	Jukes-Cantor	[Basefreq: =0.25]
44	1	Jukes-Cantor	[Basefreq: =0.25]
45	1	Jukes-Cantor	[Basefreq: =0.25]
46	1	Jukes-Cantor	[Basefreq: =0.25]
47	1	Jukes-Cantor	[Basefreq: =0.25]
48	1	Jukes-Cantor	[Basefreq: =0.25]
49	1	Jukes-Cantor	[Basefreq: =0.25]
50	1	Jukes-Cantor	[Basefreq: =0.25]
51	1	Jukes-Cantor	[Basefreq: =0.25]
52	1	Jukes-Cantor	[Basefreq: =0.25]
53	1	Jukes-Cantor	[Basefreq: =0.25]
54	1	Jukes-Cantor	[Basefreq: =0.25]
55	1	Jukes-Cantor	[Basefreq: =0.25]
56	1	Jukes-Cantor	[Basefreq: =0.25]
57	1	Jukes-Cantor	[Basefreq: =0.25]
58	1	Jukes-Cantor	[Basefreq: =0.25]
59	1	Jukes-Cantor	[Basefreq: =0.25]
60	1	Jukes-Cantor	[Basefreq: =0.25]
61	1	Jukes-Cantor	[Basefreq: =0.25]
62	1	Jukes-Cantor	[Basefreq: =0.25]
63	1	Jukes-Cantor	[Basefreq: =0.25]
64	1	Jukes-Cantor	[Basefreq: =0.25]
65	1	Jukes-Cantor	[Basefreq: =0.25]
66	1	Jukes-Cantor	[Basefreq: =0.25]
67	1	Jukes-Cantor	[Basefreq: =0.25]
68	1	Jukes-Cantor	[Basefreq: =0.25]
69	1	Jukes-Cantor	[Basefreq: =0.25]
70	1	Jukes-Cantor	[Basefreq: =0.25]
71	1	Jukes-Cantor	[Basefreq: =0.25]
72	1	Jukes-Cantor	[Basefreq: =0.25]
73	1	Jukes-Cantor	[Basefreq: =0.25]
74	1	Jukes-Cantor	[Basefreq: =0.25]
75	1	Jukes-Cantor	[Basefreq: =0.25]
76	1	Jukes-Cantor	[Basefreq: =0.25]
77	1	Jukes-Cantor	[Basefreq: =0.25]
78	1	Jukes-Cantor	[Basefreq: =0.25]
79	1	Jukes-Cantor	[Basefreq: =0.25]
L			

				AUTO 5
80	1	Jukes-Cantor	[Basefreq: =0.25]	
81	1	Jukes-Cantor	[Basefreq: =0.25]	
82	1	Jukes-Cantor	[Basefreq: =0.25]	
83	1	Jukes-Cantor	[Basefreq: =0.25]	
84	1	Jukes-Cantor	[Basefreq: =0.25]	
85	1	Jukes-Cantor	[Basefreq: =0.25]	
86	1	Jukes-Cantor	[Basefreq: =0.25]	
87	1	Jukes-Cantor	[Basefreq: =0.25]	
88	1	Jukes-Cantor	[Basefreq: =0.25]	
89	1	Jukes-Cantor	[Basefreq: =0.25]	
90	1	Jukes-Cantor	[Basefreq: =0.25]	
91	1	Jukes-Cantor	[Basefreq: =0.25]	
92	1	Jukes-Cantor	[Basefreq: =0.25]	
93	1	Jukes-Cantor	[Basefreq: =0.25]	
94	1	Jukes-Cantor	[Basefreq: =0.25]	
95	1	Jukes-Cantor	[Basefreq: =0.25]	
96	1	Jukes-Cantor	[Basefreq: =0.25]	
97	1	Jukes-Cantor	[Basefreq: =0.25]	
98	1	Jukes-Cantor	[Basefreq: =0.25]	
99	1	Jukes-Cantor	[Basefreq: =0.25]	
100	1	Jukes-Cantor	[Basefreq: =0.25]	
Sites per	locus			
Locus		Sites		
1	1	0000		

Locus	Sites
1	10000
2	10000
3	10000
4	10000
5	10000
6	10000
7	10000
8	10000
9	10000
10	10000
11	10000
12	10000
13	10000
14	10000
15	10000
16	10000
17	10000
18	10000
19	10000
20	10000

		, in the second
21	10000	
22	10000	
23	10000	
24	10000	
25	10000	
26	10000	
27	10000	
28	10000	
29	10000	
30	10000	
31	10000	
32	10000	
33	10000	
34	10000	
35	10000	
36	10000	
37	10000	
38	10000	
39	10000	
40	10000	
41	10000	
42	10000	
43	10000	
44	10000	
45	10000	
46	10000	
47	10000	
48	10000	
49	10000	
50	10000	
51	10000	
52	10000	
53	10000	
54	10000	
55	10000	
56	10000	
57	10000	
58	10000	
59	10000	
60	10000	
61	10000	
62	10000	
63	10000	
64	10000	
65	10000	
l		

66	10000				
67	10000				
68	10000				
69	10000				
70	10000				
71	10000				
72	10000				
73	10000				
74	10000				
75	10000				
76	10000				
77	10000				
78	10000				
79	10000				
80	10000				
81	10000				
82	10000				
83	10000				
84	10000				
85	10000				
86	10000				
87	10000				
88	10000				
89	10000				
90	10000				
91	10000				
92	10000				
93	10000				
94	10000				
95	10000				
96	10000				
97	10000				
98	10000				
99	10000				
100	10000				
Site rate	e variation and probat	oilities:			
Locus S	Sublocus Region type	Rate of change	Probability	Patch size	
1	1 1	1.000	1.000	1.000	
2	1 1	1.000	1.000	1.000	
3	1 1	1.000	1.000	1.000	
4	1 1	1.000	1.000	1.000	
5	1 1	1.000	1.000	1.000	
6	1 1	1.000	1.000	1.000	

7 8 9 10 11	1 1 1 1	1 1 1	1.000 1.000	1.000 1.000	1.000 1.000	
9 10	1		1.000	1 ()()()	1 ()()()	
10			1.000	1.000	1.000	
		1	1.000	1.000	1.000	
	1	1	1.000	1.000	1.000	
12	1	1	1.000	1.000	1.000	
13	1	1	1.000	1.000	1.000	
14	1	1	1.000	1.000	1.000	
15	1	1	1.000	1.000	1.000	
16	1	1	1.000	1.000	1.000	
17	1	1	1.000	1.000	1.000	
18	1	1	1.000	1.000	1.000	
19	1	1	1.000	1.000	1.000	
20	1	1	1.000	1.000	1.000	
21	1	1	1.000	1.000	1.000	
22	1	1	1.000	1.000	1.000	
23	1	1	1.000	1.000	1.000	
24	1	1	1.000	1.000	1.000	
25	1	1	1.000	1.000	1.000	
26	1	1	1.000	1.000	1.000	
27	1	1	1.000	1.000	1.000	
28	1	1	1.000	1.000	1.000	
29	1	1	1.000	1.000	1.000	
30	1	1	1.000	1.000	1.000	
31	1	1	1.000	1.000	1.000	
32	1	1	1.000	1.000	1.000	
33	1	1	1.000	1.000	1.000	
34	1	1	1.000	1.000	1.000	
35	1	1	1.000	1.000	1.000	
36	1	1	1.000	1.000	1.000	
37	1	1	1.000	1.000	1.000	
38	1	1	1.000	1.000	1.000	
39	1	1	1.000	1.000	1.000	
40	1	1	1.000	1.000	1.000	
41	1	1	1.000	1.000	1.000	
42	1	1	1.000	1.000	1.000	
43	1	1	1.000	1.000	1.000	
44	1	1	1.000	1.000	1.000	
45	1	1	1.000	1.000	1.000	
46	1	1	1.000	1.000	1.000	
47	1	1	1.000	1.000	1.000	
48	1	1	1.000	1.000	1.000	
49	1	1	1.000	1.000	1.000	
50	1	1	1.000	1.000	1.000	
51	1	1	1.000	1.000	1.000	

52	1	1	1.000	1.000	1.000	
53	1	1	1.000	1.000	1.000	
54	1	1	1.000	1.000	1.000	
55	1	1	1.000	1.000	1.000	
56	1	1	1.000	1.000	1.000	
57	1	1	1.000	1.000	1.000	
58	1	1	1.000	1.000	1.000	
59	1	1	1.000	1.000	1.000	
60	1	1	1.000	1.000	1.000	
61	1	1	1.000	1.000	1.000	
62	1	1	1.000	1.000	1.000	
63	1	1	1.000	1.000	1.000	
64	1	1	1.000	1.000	1.000	
65	1	1	1.000	1.000	1.000	
66	1	1	1.000	1.000	1.000	
67	1	1	1.000	1.000	1.000	
68	1	1	1.000	1.000	1.000	
69	1	1	1.000	1.000	1.000	
70	1	1	1.000	1.000	1.000	
71	1	1	1.000	1.000	1.000	
72	1	1	1.000	1.000	1.000	
73	1	1	1.000	1.000	1.000	
74	1	1	1.000	1.000	1.000	
75	1	1	1.000	1.000	1.000	
76	1	1	1.000	1.000	1.000	
77	1	1	1.000	1.000	1.000	
78	1	1	1.000	1.000	1.000	
79	1	1	1.000	1.000	1.000	
80	1	1	1.000	1.000	1.000	
81	1	1	1.000	1.000	1.000	
82	1	1	1.000	1.000	1.000	
83	1	1	1.000	1.000	1.000	
84	1	1	1.000	1.000	1.000	
85	1	1	1.000	1.000	1.000	
86	1	1	1.000	1.000	1.000	
87	1	1	1.000	1.000	1.000	
88	1	1	1.000	1.000	1.000	
89	1	1	1.000	1.000	1.000	
90	1	1	1.000	1.000	1.000	
91	1	1	1.000	1.000	1.000	
92	1	1	1.000	1.000	1.000	
93	1	1	1.000	1.000	1.000	
94	1	1	1.000	1.000	1.000	
95	1	1	1.000	1.000	1.000	
96	1	1	1.000	1.000	1.000	
L						

97		1	1.000	1.000	1.000	
98	1 1	1	1.000	1.000	1.000	
99	1	1	1.000	1.000	1.000	
100	1	1	1.000	1.000	1.000	
Population		•	1.000	1.000	Locus	Gene copies
1 Romans					1	10
1 Roman	5110111_0				2	10
					3	10
					4	10
					5	10
					6	10
					7	10
					8	10
					9	10
					10	10
					11	10
					12	10
					13	10
					14	10
					15	10
					16	10
					17	10
					18	10
					19	10
					20	10
					21	10
					22	10
					23	10
					24	10
					25	10
					26	10
					27	10
					28	10
					29	10
					30	10
					31	10
					32	10
					33	10
					33 34	
					34 35	10
						10
					36	10
					37	10
					38	10
					39	10
					40	10

41	10
42	10
43	10
44	10
45	10
46	10
47	10
48	10
49	10
50	10
51	10
52	10
53	10
54	10
55	10
56	10
57	10
58	10
59	10
60	10
61	10
62	10
63	10
64	10
65	10
66	10
67	10
68	10
69	10
70	10
71	10
72	10
73	10
74	10
75	10
76	10
77	10
78	10
79	10
80	10
81	10
82	10
83	10
84	10
85	10

	86	10	
	87	10	
	88	10	
	89	10	
	90	10	
	91	10	
	92	10	
	93	10	
	94	10	
	95	10	
	96	10	
	97	10	
	98	10	
	99	10	
	100	10	
Total of all populations	1	10	
	2	10	
	3	10	
	4	10	
	5	10	
	6	10	
	7	10	
	8	10	
	9	10	
	10	10	
	11	10	
	12	10	
	13	10	
	14	10	
	15	10	
	16	10	
	17	10	
	18	10	
	19	10	
	20	10	
	21	10	
	22	10	
	23	10	
	24	10	
	25	10	
	26	10	
	27	10	
	28	10	
	29	10	
	30	10	

31	10
32	10
33	10
34	10
35	10
36	10
37	10
38	10
39	10
40	10
41	10
42	10
43	10
44	10
45	10
46	10
47	10
48	10
49	10
50	10
51	10
52	10
53	10
54	10
55	10
56	10
57	10
58	10
59	10
60	10
61	10
62	10
63	10
64	10
65	10
66	10
67	10
68	10
69	10
70	10
71	10
72	10
73	10
74	10
75	10

76	10
77	10
78	10
79	10
80	10
81	10
82	10
83	10
84	10
85	10
86	10
87	10
88	10
89	10
90	10
91	
	10
92	10
93	10
94	10
95	10
96	10
97	10
98	10
99	10
100	10

# Bayesian Analysis: Posterior distribution table

Locus	Parameter	2.5%	25.0%	Mode	75.0%	97.5%	Median	Mean
1	$\Theta_1$	0.03113	0.04440	0.04783	0.04960	0.05153	0.04457	0.08151
2	$\Theta_1$	0.03053	0.04353	0.04783	0.04973	0.05167	0.04470	0.08324
3	$\Theta_1$	0.03380	0.04400	0.04783	0.04967	0.05153	0.04523	0.08370
4	$\Theta_1$	0.03453	0.04580	0.04803	0.04973	0.05173	0.04597	0.08668
5	$\Theta_1$	0.03247	0.04373	0.04790	0.04973	0.05167	0.04497	0.08254
6	$\Theta_1$	0.03460	0.04573	0.04777	0.04913	0.05160	0.04597	0.08730
7	$\Theta_1$	0.03280	0.04400	0.04777	0.04980	0.05167	0.04510	0.08429
8	$\Theta_1$	0.03427	0.04460	0.04790	0.04980	0.05173	0.04577	0.08538
9	$\Theta_1$	0.03213	0.04353	0.04777	0.04953	0.05167	0.04477	0.08434
10	$\Theta_1$	0.03353	0.04440	0.04790	0.04973	0.05160	0.04557	0.08472
11	$\Theta_1$	0.03093	0.04340	0.04783	0.04980	0.05153	0.04457	0.08177
12	$\Theta_1$	0.03207	0.04393	0.04790	0.04987	0.05160	0.04510	0.08302
13	$\Theta_1$	0.03033	0.04313	0.04783	0.04980	0.05153	0.04437	0.08062
14	$\Theta_1$	0.03340	0.04420	0.04797	0.04980	0.05167	0.04543	0.08506
15	$\Theta_1$	0.03267	0.04513	0.04797	0.04953	0.05167	0.04543	0.08547
16	$\Theta_1$	0.03373	0.04547	0.04797	0.04960	0.05153	0.04563	0.08546
17	$\Theta_1$	0.03273	0.04407	0.04790	0.04967	0.05167	0.04530	0.08542
18	$\Theta_1$	0.03033	0.04393	0.04790	0.04953	0.05153	0.04410	0.08003

Migrate 5.0.0a: (http://popgen.sc.fsu.edu) [program run on 19:59:31]

19	$\Theta_1$	0.02667	0.04220	0.04763	0.04940	0.05140	0.04257	0.07494
20	$\Theta_1$	0.03167	0.04380	0.04797	0.04987	0.05160	0.04497	0.08313
21	$\Theta_1$	0.03160	0.04360	0.04790	0.04980	0.05160	0.04477	0.08187
22	$\Theta_1$	0.03367	0.04460	0.04790	0.04987	0.05160	0.04577	0.08545
23	$\Theta_1$	0.03387	0.04427	0.04803	0.04973	0.05173	0.04550	0.08545
24	$\Theta_1$	0.03413	0.04460	0.04790	0.04987	0.05167	0.04570	0.08535
25	$\Theta_1$	0.03200	0.04467	0.04790	0.04953	0.05160	0.04483	0.08316
26	$\Theta_1$	0.03280	0.04500	0.04783	0.04933	0.05160	0.04517	0.08435
27	$\Theta_1$	0.03107	0.04313	0.04790	0.04953	0.05160	0.04443	0.08108
28	$\Theta_1$	0.03280	0.04473	0.04783	0.04940	0.05153	0.04490	0.08256
29	$\Theta_1$	0.02967	0.04280	0.04783	0.04973	0.05153	0.04403	0.08079
30	$\Theta_1$	0.03427	0.04480	0.04810	0.05007	0.05167	0.04583	0.08604
31	$\Theta_1$	0.03247	0.04413	0.04797	0.04987	0.05167	0.04530	0.08433
32	$\Theta_1$	0.03233	0.04387	0.04790	0.04980	0.05153	0.04503	0.08275
33	$\Theta_1$	0.03393	0.04520	0.04790	0.04967	0.05173	0.04577	0.08569
34	$\Theta_1$	0.03287	0.04433	0.04790	0.04980	0.05153	0.04543	0.08471
35	$\Theta_1$	0.03107	0.04367	0.04783	0.04987	0.05167	0.04483	0.08257
36	$\Theta_1$	0.03420	0.04513	0.04790	0.04967	0.05167	0.04530	0.08655
37	$\Theta_1$	0.03467	0.04480	0.04803	0.05000	0.05153	0.04583	0.08713
38	$\Theta_1$	0.03327	0.04393	0.04790	0.04967	0.05153	0.04517	0.08543
39	$\Theta_1$	0.03260	0.04387	0.04783	0.04973	0.05160	0.04510	0.08404
40	$\Theta_1$	0.03173	0.04413	0.04770	0.04967	0.05160	0.04510	0.08403
41	$\Theta_1$	0.02967	0.04433	0.04790	0.04960	0.05160	0.04450	0.08282

_ocus	Parameter	2.5%	25.0%	Mode	75.0%	97.5%	Median	Mean
42	$\Theta_1$	0.03160	0.04360	0.04777	0.04967	0.05160	0.04483	0.08375
43	$\Theta_1$	0.03273	0.04380	0.04803	0.05007	0.05167	0.04530	0.08428
44	$\Theta_1$	0.03313	0.04473	0.04803	0.05000	0.05173	0.04583	0.08647
45	$\Theta_1$	0.03147	0.04333	0.04777	0.04967	0.05153	0.04457	0.08165
46	$\Theta_1$	0.03127	0.03493	0.04797	0.05133	0.05167	0.04497	0.08324
47	$\Theta_1$	0.03320	0.04413	0.04797	0.04973	0.05160	0.04537	0.08537
48	$\Theta_1$	0.03167	0.04360	0.04783	0.04980	0.05160	0.04477	0.08166
49	$\Theta_1$	0.03373	0.04473	0.04797	0.04987	0.05160	0.04583	0.08627
50	$\Theta_1$	0.03193	0.04360	0.04783	0.04967	0.05167	0.04483	0.08178
51	$\Theta_1$	0.03347	0.04447	0.04777	0.04973	0.05153	0.04557	0.08516
52	$\Theta_1$	0.03307	0.04420	0.04797	0.04993	0.05153	0.04530	0.08342
53	$\Theta_1$	0.03213	0.03900	0.04783	0.05067	0.05160	0.04497	0.08504
54	$\Theta_1$	0.03247	0.04380	0.04770	0.04940	0.05153	0.04517	0.08489
55	$\Theta_1$	0.03053	0.04327	0.04777	0.04973	0.05160	0.04443	0.08098
56	$\Theta_1$	0.03153	0.03780	0.04777	0.05087	0.05153	0.04490	0.08300
57	$\Theta_1$	0.03327	0.04500	0.04777	0.04913	0.05153	0.04517	0.08432
58	$\Theta_1$	0.03213	0.04413	0.04777	0.04967	0.05153	0.04530	0.08406
59	$\Theta_1$	0.03240	0.04460	0.04803	0.04973	0.05180	0.04537	0.08384
60	$\Theta_1$	0.03087	0.04380	0.04797	0.04993	0.05160	0.04490	0.08436
61	$\Theta_1$	0.03093	0.04353	0.04790	0.04980	0.05160	0.04470	0.08277

62	$\Theta_1$	0.03413	0.04480	0.04803	0.05000	0.05160	0.04583	0.08650
63	$\Theta_1$	0.03373	0.04447	0.04783	0.04973	0.05160	0.04570	0.08446
64	$\Theta_1$	0.03413	0.04440	0.04810	0.05007	0.05167	0.04597	0.08657
65	$\Theta_1$	0.03253	0.04393	0.04777	0.04973	0.05153	0.04510	0.08389
66	$\Theta_1$	0.03360	0.04513	0.04783	0.04933	0.05153	0.04530	0.08419
67	$\Theta_1$	0.02973	0.04367	0.04770	0.04973	0.05167	0.04483	0.08356
68	$\Theta_1$	0.03520	0.04487	0.04803	0.04987	0.05160	0.04603	0.08728
69	$\Theta_1$	0.03220	0.04400	0.04783	0.04973	0.05167	0.04517	0.08441
70	$\Theta_1$	0.03320	0.04453	0.04817	0.04993	0.05180	0.04570	0.08697
71	$\Theta_1$	0.03333	0.04420	0.04790	0.04987	0.05160	0.04530	0.08440
72	$\Theta_1$	0.02593	0.04247	0.04777	0.04947	0.05187	0.04383	0.07850
73	$\Theta_1$	0.03087	0.04393	0.04777	0.04947	0.05153	0.04410	0.08040
74	$\Theta_1$	0.03113	0.04347	0.04783	0.04973	0.05153	0.04463	0.08264
75	$\Theta_1$	0.03467	0.04453	0.04777	0.04953	0.05167	0.04583	0.08564
76	$\Theta_1$	0.03040	0.04393	0.04777	0.04947	0.05153	0.04410	0.07917
77	$\Theta_1$	0.03047	0.04320	0.04783	0.04973	0.05153	0.04437	0.08230
78	$\Theta_1$	0.03353	0.04547	0.04810	0.04973	0.05167	0.04563	0.08586
79	$\Theta_1$	0.03387	0.04447	0.04763	0.04960	0.05173	0.04570	0.08601
80	$\Theta_1$	0.03087	0.04320	0.04783	0.04973	0.05160	0.04443	0.08107
81	$\Theta_1$	0.03253	0.04420	0.04783	0.04987	0.05153	0.04530	0.08402
82	$\Theta_1$	0.03280	0.03693	0.04797	0.05120	0.05160	0.04530	0.08466
83	$\Theta_1$	0.03473	0.04487	0.04797	0.04987	0.05173	0.04597	0.08702
84	$\Theta_1$	0.03413	0.04467	0.04803	0.04993	0.05167	0.04577	0.08721

_ocus	Parameter	2.5%	25.0%	Mode	75.0%	97.5%	Median	Mean
85	$\Theta_1$	0.03380	0.04460	0.04777	0.04967	0.05160	0.04570	0.08562
86	$\Theta_1$	0.03213	0.04420	0.04797	0.04987	0.05160	0.04537	0.08489
87	$\Theta_1$	0.03320	0.04440	0.04790	0.04973	0.05160	0.04557	0.08373
88	$\Theta_1$	0.03313	0.04473	0.04790	0.04987	0.05160	0.04583	0.08598
89	$\Theta_1$	0.03340	0.04420	0.04790	0.04973	0.05153	0.04537	0.08544
90	$\Theta_1$	0.03307	0.04400	0.04783	0.04960	0.05167	0.04523	0.08463
91	$\Theta_1$	0.03213	0.04340	0.04797	0.05000	0.05160	0.04517	0.08332
92	$\Theta_1$	0.03427	0.04527	0.04797	0.04953	0.05160	0.04543	0.08697
93	$\Theta_1$	0.03127	0.04307	0.04783	0.04953	0.05147	0.04443	0.08199
94	$\Theta_1$	0.03387	0.04413	0.04803	0.04953	0.05153	0.04517	0.08599
95	$\Theta_1$	0.03400	0.04433	0.04797	0.04980	0.05167	0.04550	0.08604
96	$\Theta_1$	0.03187	0.04367	0.04783	0.04973	0.05153	0.04483	0.08230
97	$\Theta_1$	0.03527	0.04500	0.04797	0.04987	0.05180	0.04610	0.08716
98	$\Theta_1$	0.03233	0.04433	0.04797	0.05000	0.05173	0.04543	0.08486
99	$\Theta_1$	0.03393	0.04447	0.04803	0.04987	0.05173	0.04557	0.08560
100	$\Theta_1$	0.03207	0.04400	0.04777	0.04973	0.05160	0.04517	0.08444
All	$\Theta_1$	0.00800	0.01340	0.01503	0.01673	0.01933	0.01483	0.09964

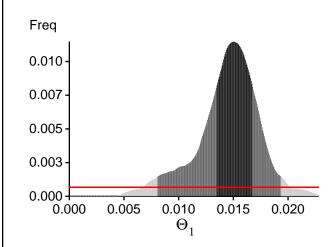
#### Citation suggestions:

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



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

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

ocus.	TI(1a)	BTI(1b)	SS(2)	HS(3)
1	-15175.34	-14776.67	-14813.21	-14874.82
2	-14930.09	-14616.85	-14669.92	-14729.13
3	-14769.40	-14457.03	-14505.03	-14567.54
4	-18414.44	-16380.14	-16122.19	-16181.61
5	-14880.18	-14443.54	-14466.64	-14534.63
6	-17643.07	-16358.05	-16253.65	-16302.28
7	-15624.01	-15034.12	-15040.36	-15098.72
8	-15765.71	-15102.79	-15094.94	-15153.05
9	-15117.70	-14782.01	-14837.42	-14891.94
10	-15110.57	-14637.92	-14662.40	-14720.28
11	-21859.67	-18199.83	-17645.90	-17710.49
12	-15086.77	-14650.61	-14679.76	-14739.70
13	-14471.06	-14221.28	-14274.84	-14341.86
14	-15051.77	-14621.10	-14654.79	-14711.45
15	-18087.51	-16354.53	-16155.01	-16213.39
16	-15160.41	-14852.86	-14913.29	-14965.60
17	-16002.72	-15351.98	-15352.08	-15409.82
18	-14396.47	-14181.23	-14238.58	-14308.18
19	-14218.83	-14051.16	-14107.01	-14183.91
20	-14871.21	-14513.48	-14555.71	-14618.35
21	-15612.66	-14898.05	-14872.17	-14937.12
22	-15190.63	-14736.17	-14766.75	-14826.76
23	-16714.82	-15921.60	-15898.98	-15953.73
24	-21831.34	-18319.77	-17793.26	-17853.48
25	-14992.45	-14647.71	-14695.83	-14754.18
26	-15243.08	-14789.89	-14817.94	-14876.72
27	-15903.96	-15228.79	-15216.78	-15281.20
28	-15729.24	-15079.47	-15069.80	-15133.83
29	-14401.99	-14172.15	-14230.14	-14296.05

Migrate 5.0.0a: (http://popgen.sc.fsu.edu) [program run on 19:59:31]

30	-18528.16	-16427.40	-16158.23	-16220.65
31	-16368.79	-15823.92	-15846.41	-15904.89
32	-14659.07	-14374.34	-14425.52	-14491.22
33	-16894.31	-15991.29	-15944.07	-15999.91
34	-15484.01	-14944.78	-14959.51	-15017.35
35	-15335.69	-14870.89	-14893.56	-14955.70
36	-20266.10	-18713.99	-18585.03	-18637.70
37	-18392.08	-16552.82	-16336.50	-16393.98
38	-15963.89	-15301.58	-15295.41	-15349.90
39	-14952.69	-14528.95	-14557.94	-14617.90
40	-14907.40	-14565.40	-14612.11	-14672.52
41	-20457.74	-17916.55	-17579.45	-17639.04
42	-15249.08	-14838.80	-14877.85	-14936.41
43	-15219.79	-14782.46	-14812.90	-14872.94
44	-15300.94	-14894.27	-14938.12	-14991.13
45	-14764.18	-14408.61	-14449.00	-14513.46
46	-15106.68	-14681.44	-14713.16	-14772.45
47	-15360.22	-14812.56	-14816.98	-14880.99
48	-15642.34	-14920.09	-14892.81	-14958.42
49	-15579.34	-15050.82	-15071.38	-15126.68
50	-14681.52	-14381.88	-14430.38	-14495.40
51	-15798.54	-15117.18	-15106.56	-15163.78
52	-15901.29	-15063.68	-15019.46	-15082.45
53	-15196.25	-14811.54	-14855.15	-14912.58
54	-15546.68	-15001.77	-15014.38	-15075.19
55	-14614.83	-14317.78	-14364.64	-14430.90
56	-14952.55	-14611.18	-14657.05	-14716.65
57	-15641.90	-15202.58	-15235.90	-15297.87
58	-15825.63	-15075.24	-15050.84	-15110.27
59	-15269.85	-14766.32	-14773.55	-14843.59
60	-17254.37	-16311.23	-16269.13	-16322.55
61	-15783.87	-15028.40	-15000.02	-15062.24
62	-15933.02	-15366.61	-15383.23	-15438.62
63	-15114.03	-14643.21	-14666.11	-14726.28
64	-17161.50	-16095.82	-16019.94	-16077.98
65	-16580.56	-15671.70	-15619.18	-15684.71
66	-14880.97	-14606.08	-14661.28	-14722.99
67	-15479.18	-15014.70	-15037.38	-15100.75
68	-15848.91	-15387.70	-15423.21	-15480.67
69	-15897.42	-15293.69	-15302.16	-15357.48
70	-17759.15	-16226.66	-16069.30	-16124.67
71	-15182.35	-14720.74	-14747.30	-14806.25
72	-14266.75	-14057.71	-14116.33	-14183.82
73	-15272.68	-14922.74	-14966.79	-15032.51
74	-14616.57	-14351.99	-14405.13	-14471.20

75	-20476.45	-17345.57	-16892.13	-16948.26
76	-15514.45	-15013.60	-15032.46	-15096.19
77	-30154.85	-23649.02	-22601.91	-22668.67
78	-15580.89	-14974.12	-14977.18	-15034.69
79	-15353.14	-14934.29	-14978.49	-15031.07
80	-15722.68	-15200.64	-15219.24	-15282.80
81	-14996.41	-14681.48	-14732.95	-14794.29
82	-15946.15	-15384.96	-15402.41	-15457.85
83	-19068.02	-17217.67	-17014.88	-17066.74
84	-16622.61	-15772.81	-15742.71	-15794.84
85	-15835.55	-15413.20	-15451.52	-15510.89
86	-15460.16	-14975.55	-15002.07	-15057.56
87	-15378.95	-14997.10	-15039.11	-15101.96
88	-18330.07	-16788.55	-16633.49	-16690.08
89	-15317.57	-14865.59	-14896.24	-14953.05
90	-15907.56	-15330.30	-15339.36	-15396.52
91	-15237.60	-14808.12	-14842.02	-14899.58
92	-16912.45	-15711.72	-15612.07	-15664.42
93	-15124.48	-14764.28	-14802.53	-14865.56
94	-15849.24	-15240.74	-15251.71	-15303.99
95	-18060.14	-16871.52	-16785.76	-16838.52
96	-14974.07	-14572.18	-14606.15	-14667.76
97	-17069.89	-15825.89	-15719.99	-15772.14
98	-14843.00	-14531.13	-14585.00	-14641.31
99	-15160.45	-14799.51	-14846.72	-14904.73
100	-14885.13	-14536.54	-14584.40	-14644.49
All	-1610789.23	-1531893.25	-1528814.46	-1534789.68

- (1a) TI: Thermodynamic integration: log(Prob(D|Model)): Good approximation with many temperatures(1b) BTI: Bezier-approximated Thermodynamic integration: when using few temperatures USE THIS!
- (2) SS: Steppingstone Sampling (Xie et al 2011)
- (3) HS: Harmonic mean approximation: Overestimates the marginal likelihood, poor variance [Scaling factor = 142.725660]

#### Citation suggestions:

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

Palczewski M. and P. Beerli, 2014. Population model comparison using multi-locus datasets. In M.-H. Chen, L. Kuo, and P. O. Lewis, editors, Bayesian Phylogenetics: Methods,

Algorithms, and Applications, pages 187-200. CRC Press, 2014.

Xie W., P. O. Lewis, Y. Fan, L. Kuo, and M.-H. Chen. 2011. Improving marginal likelihood estimation for Bayesian phylogenetic model selection. Systematic Biology, 60(2):150â 160, 2011.

## Acceptance ratios for all parameters and the genealogies

Parameter	Accepted changes	Ratio
$\Theta_1$ Genealogies	375942224/399965520 105526033/1600034480	0.93994 0.06595

## MCMC-Autocorrelation and Effective MCMC Sample Size

Parameter	Autocorrelation	Effective Sampe Size
$\Theta_1$ Genealogies	0.48763 0.38080	3455991.79 4557971.70

## Average temperatures during the run

### Chain Temperatures

- 1 0.00000
- 2 0.00000
- 3 0.00000
- 4 0.00000

Adaptive heating often fails, if the average temperatures are very close together try to rerun using static heating! If you want to compare models using marginal likelihoods then you MUST use static heating

#### Potential Problems

This section reports potential problems with your run, but such reporting is often not very accurate. Whith many parameters in a multilocus analysi s, it is very common that some parameters for some loci will not be very informative, triggering suggestions (for example to increase the prior ran ge) that are not sensible. This suggestion tool will improve with time, therefore do not blindly follow its suggestions. If some parameters are fla gged inspect the tables carefully and judge wether an action is required. For example, if you run a Bayesian

inference with sequence data, for mac roscopic 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 rou tes 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.
No warning was recorded during the run