

BEAUti 2: Standard									
File Mode View Help									
Partitions Tip Dates Site Model Clock Model Priors MCMC									
Link Site Models		Unlink Site Models		Link Clock Models		Unlink Clock Models		Link Trees	Unlink Trees
Name	File	Taxa	Sites	Data Type	Site Model	Clock Model	Tree		
COX1	COX1	11	1533	nucleotide	COX1	GTR1	linked		
ATP8	ATP8	11	156	nucleotide	GTR1	GTR1	linked		
ATP6	ATP6	11	669	nucleotide	GTR1	GTR1	linked		
COX2	COX2	11	682	nucleotide	GTR1	GTR1	linked		
COX3	COX3	11	786	nucleotide	GTR1	GTR1	linked		
CYTB	CYTB	11	1137	nucleotide	GTR1	GTR1	linked		
ND1	ND1	11	921	nucleotide	GTR2	GTR2	linked		
ND2	ND2	11	981	nucleotide	GTR1	GTR1	linked		
ND3	ND3	11	354	nucleotide	GTR1	GTR1	linked		
ND4	ND4	11	1290	nucleotide	GTR2	GTR2	linked		
ND4L	ND4L	11	291	nucleotide	GTR2	GTR2	linked		
ND5	ND5	11	1710	nucleotide	GTR2	GTR2	linked		
ND6	ND6	11	498	nucleotide	GTR1	GTR1	linked		
rrnL	rrnL	11	976	nucleotide	HKY	HKY	linked		
rrnS	rrnS	11	607	nucleotide	HKY	HKY	linked		

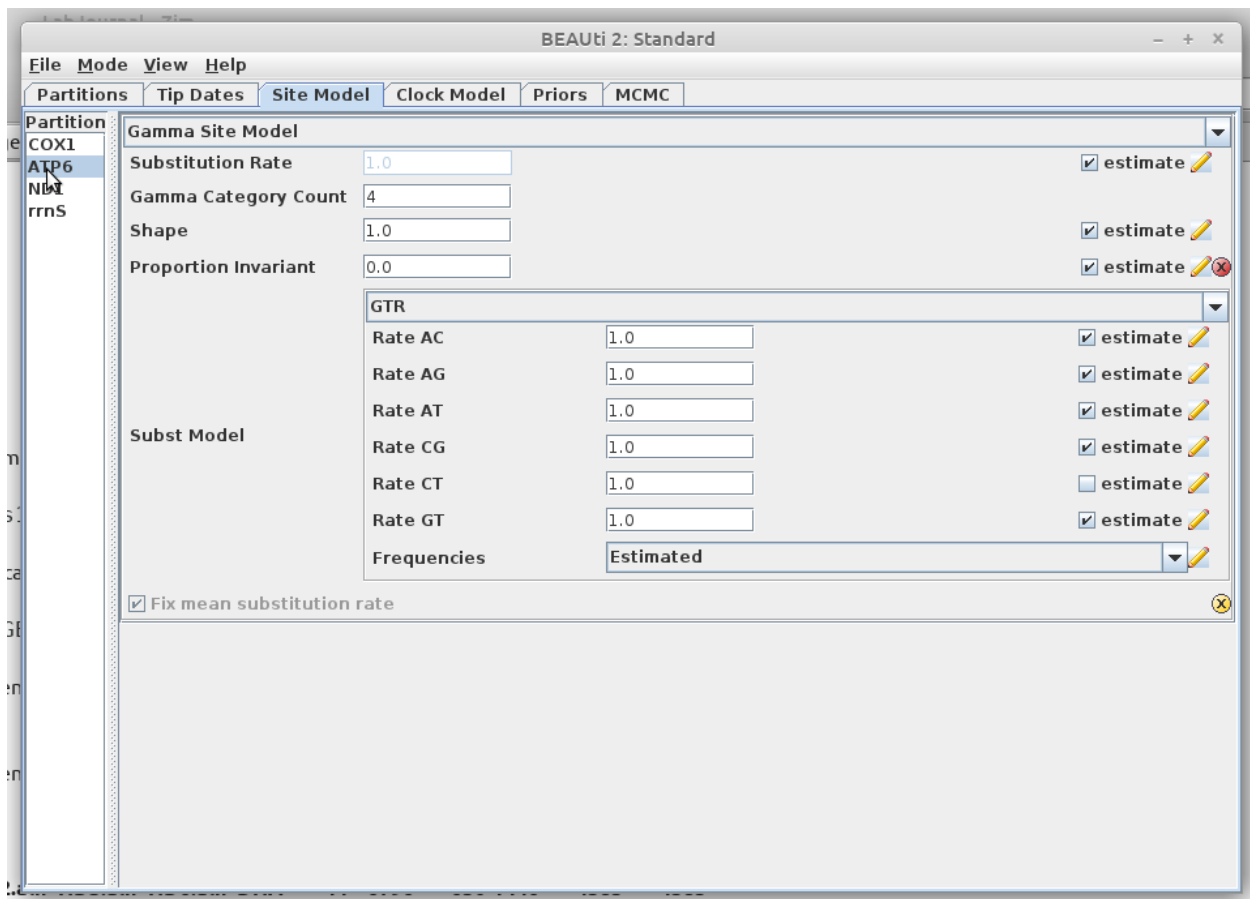
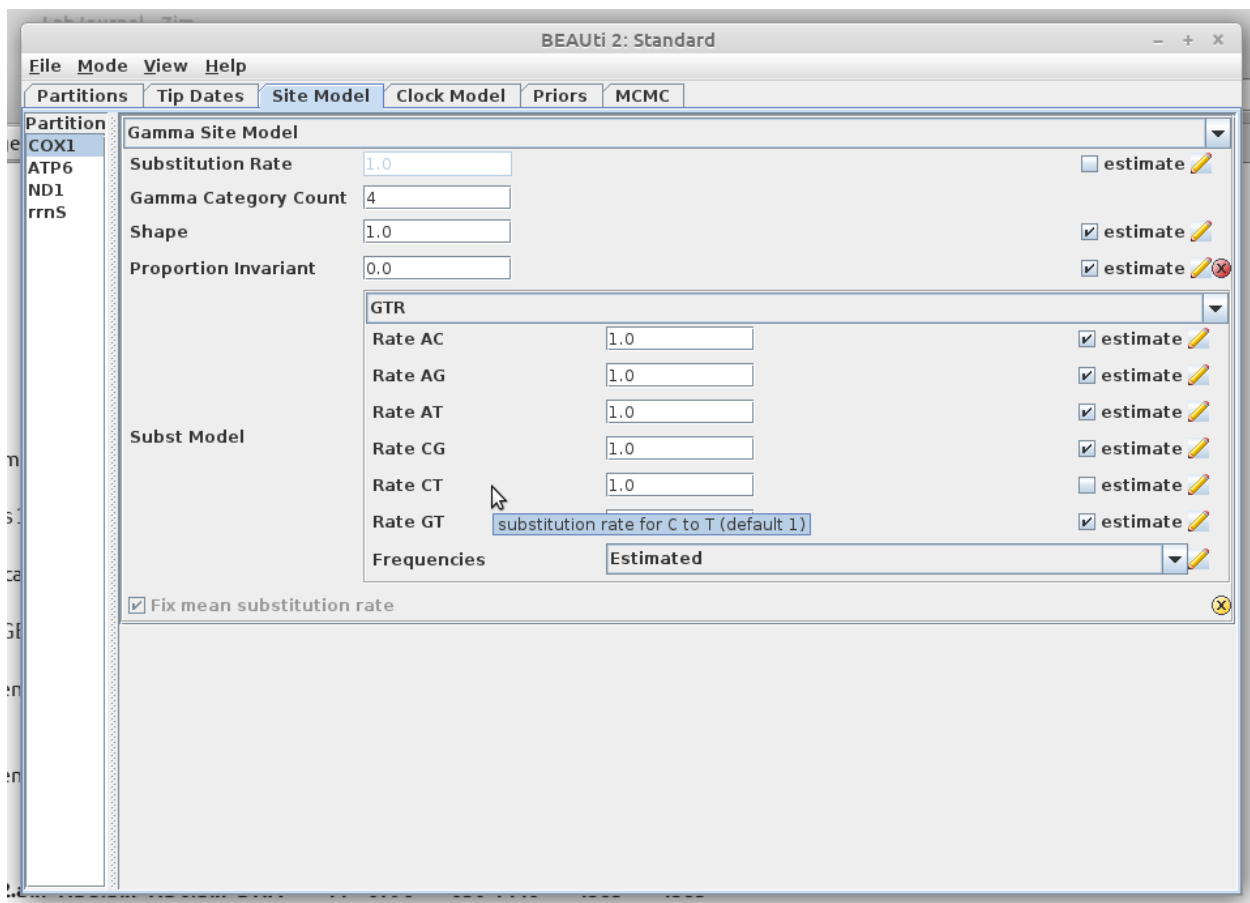
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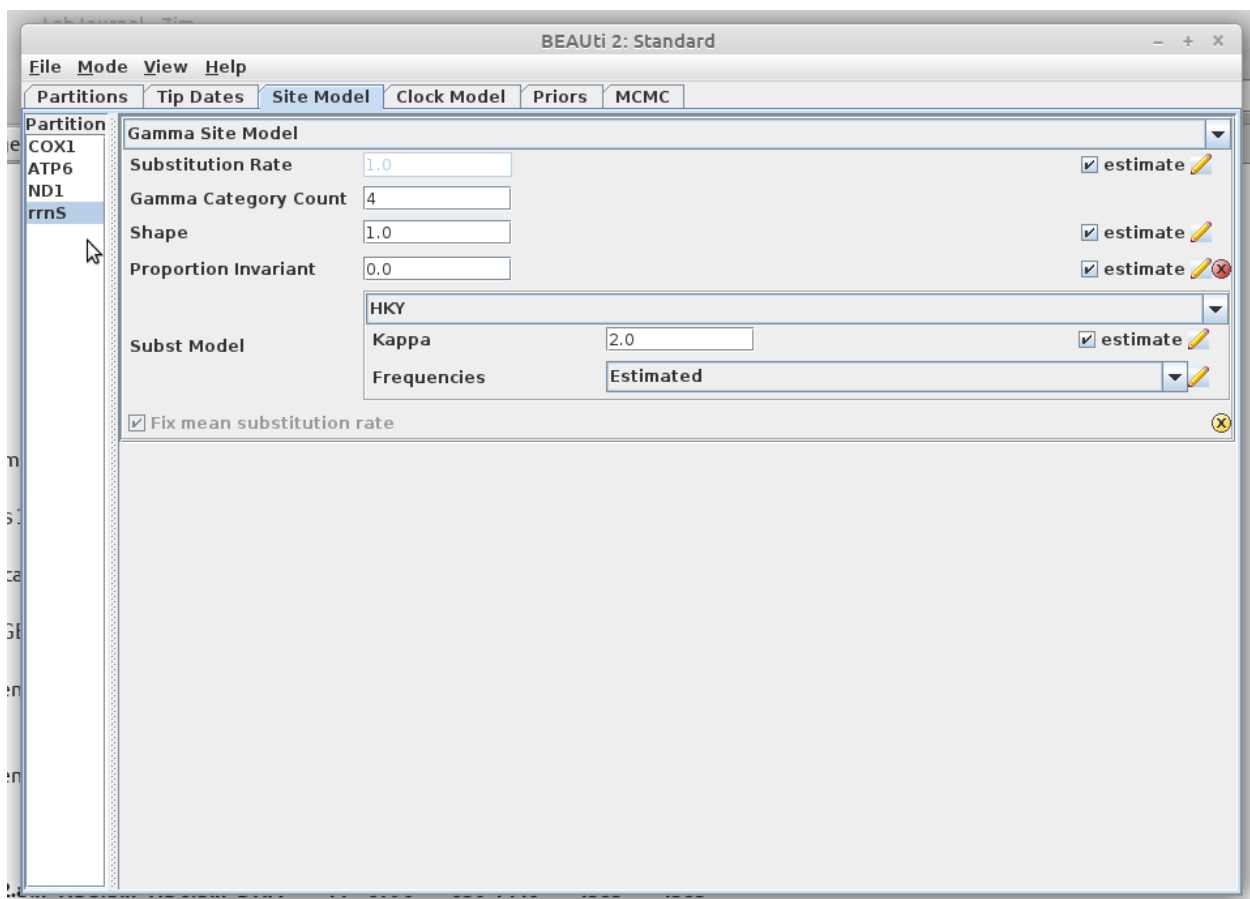
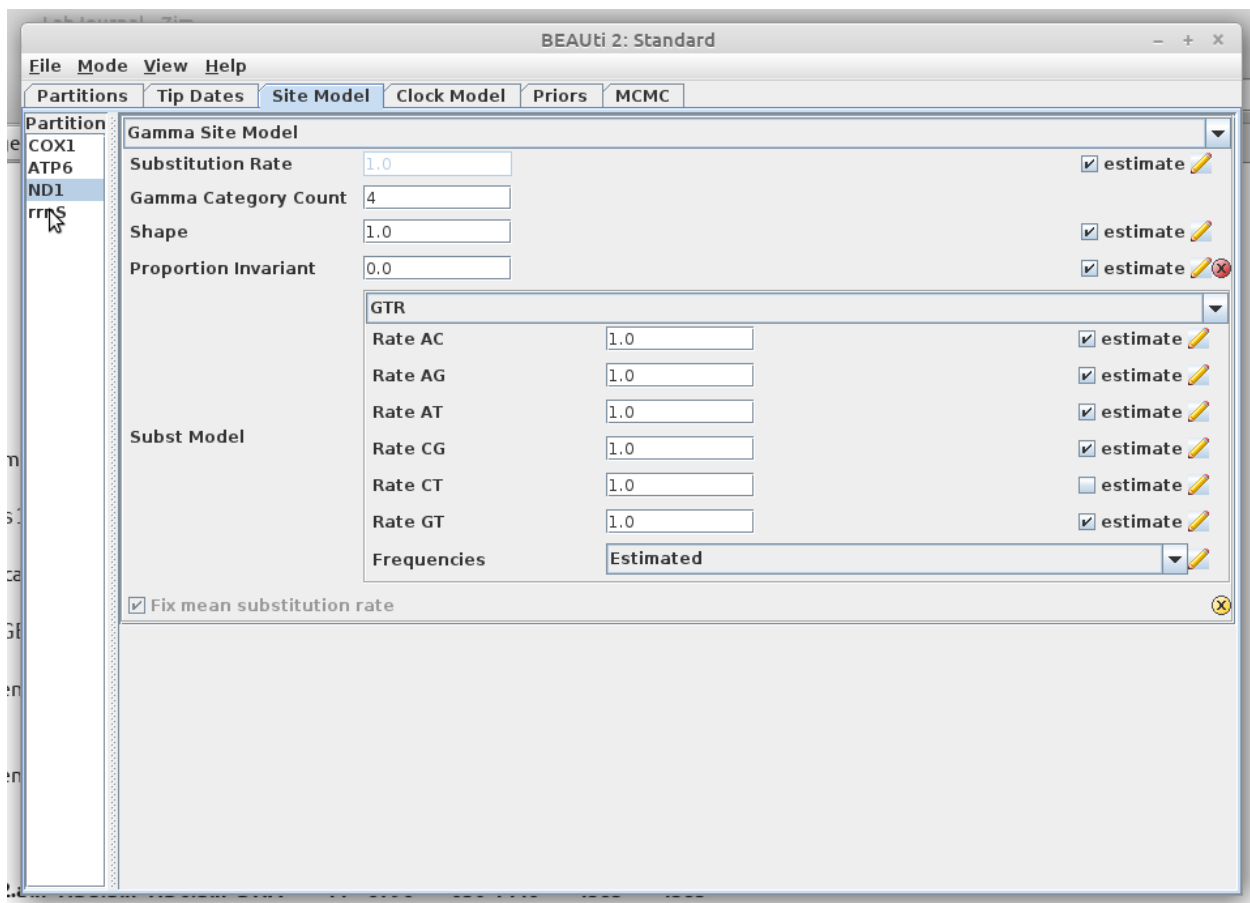
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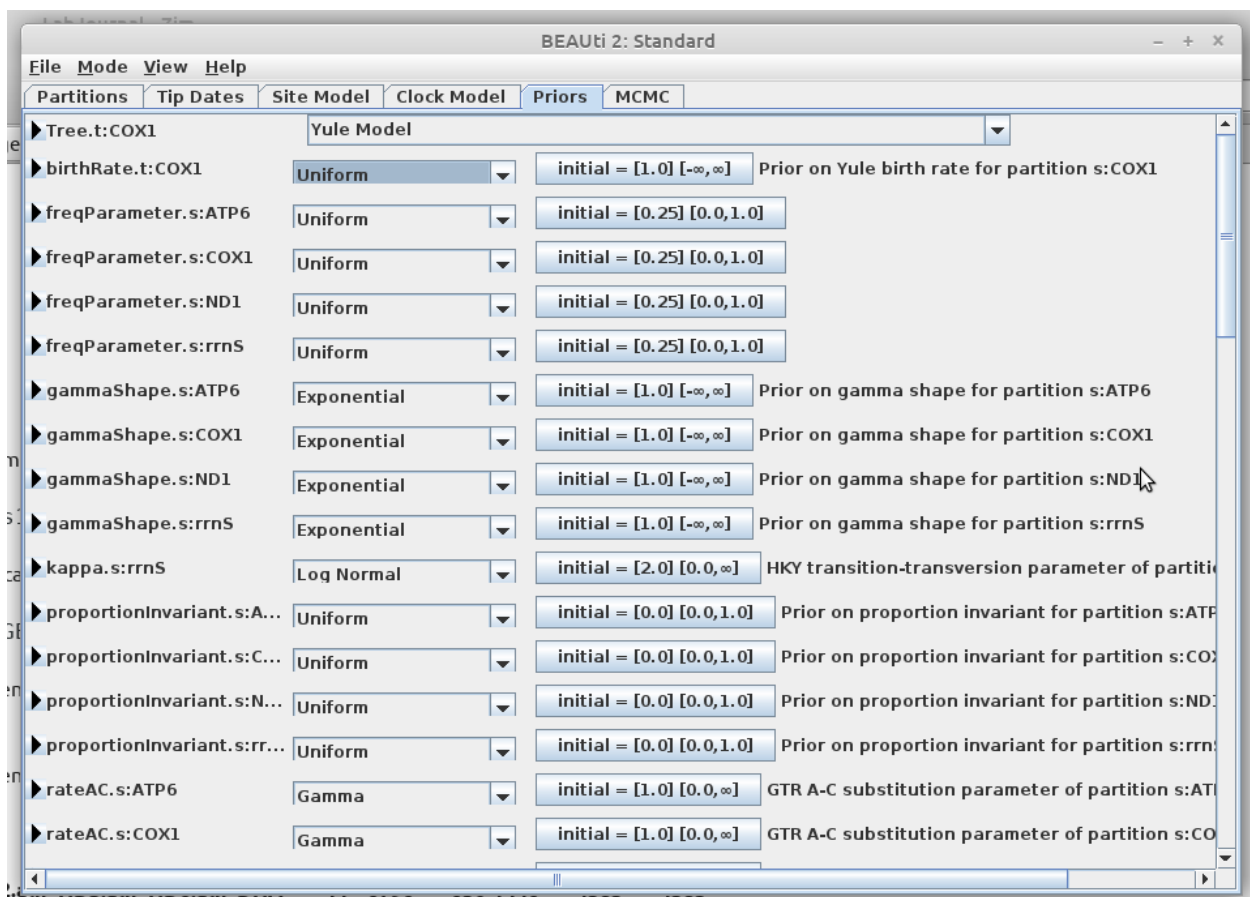
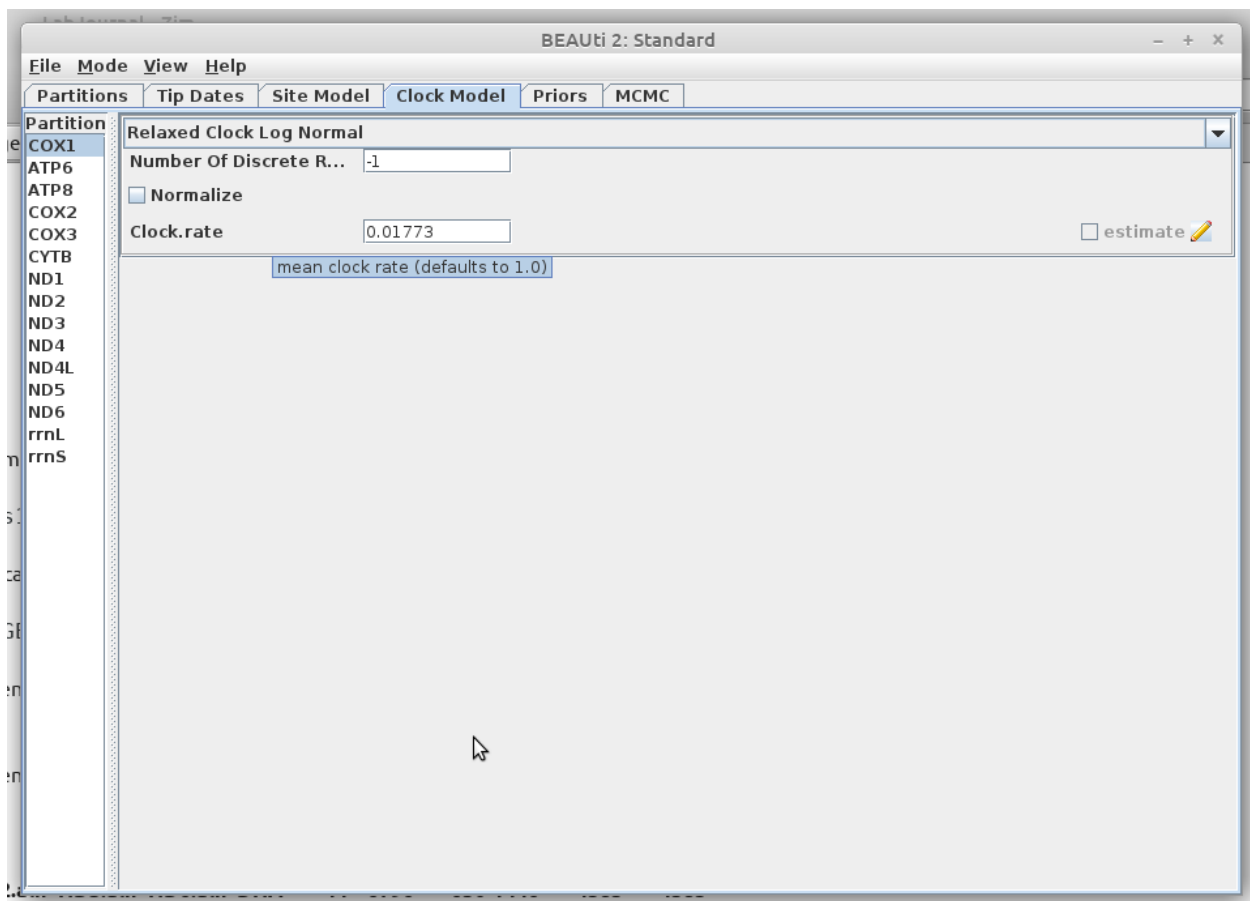
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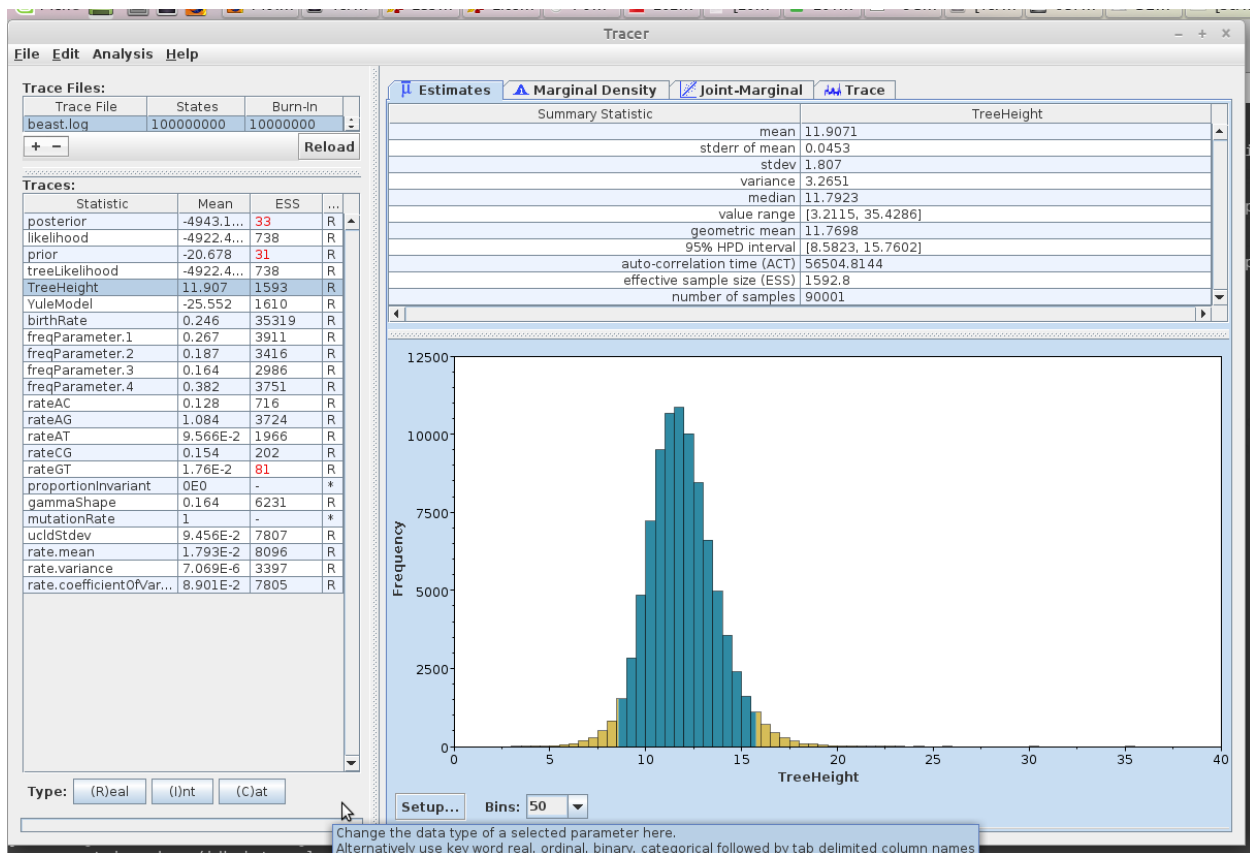
Split

BEAUti 2: Standard					
File Mode View Help					
Partitions	Tip Dates	Site Model	Clock Model	Priors	MCMC
<div> <input type="checkbox"/> Use tip dates </div>					









BEAUti 2: Standard

File Mode View Help

Partitions Tip Dates Site Model Clock Model Priors MCMC

Chain Length 10000000

Store Every -1

Pre Burnin 0

Num Initialization Atte... 10

tracelog

screenlog

File Name

Log Every 1000

Mode autodetect

Sort none

☐ Sanitise Headers

posterior

likelihood

prior

treeLog.t:COX1

☐ Sample From Prior