

Additional file 4 – Bayes factor comparisons for the selection of the clock model, the speciation process and the partition scheme

- a) Bayes factor comparisons for selection among strict and relaxed exponential and lognormal clock models under partition scheme P1: gene partition.
- b) Bayes factor comparisons for selection between Yule and Birth-death speciation models under partition scheme P1.
- c) Bayes factor comparisons for selection among partition schemes P1, P2, P3 and P1_{NOINV} (see Material and Methods for details).

HME: Harmoic mean of the likelihood from the posterior distribution value.

a)

Yule speciation process	HME	P1 strict	P1 exponential	P1 lognormal
P1 strict	-37591.196	-		
P1 exponential	-37366.070	225.126	-	
P1 lognormal	-37365.619	225.577	0.452	-

b)

Lognormal Clock	HME	P1 Yule	P1 Birth-death
P1 Yule	-37365.972	-	
P1 Birth-death	-37365.619	0.353	-

c)

Yule speciation process & Lognormal Clock	HME	P1 _{NOINV}	P1	P2	P3
P1 _{NOINV}	-39113.410	-			
P1	-37365.619	1747.792	-		
P2	-36635.311	2478.099	730.308	-	
P3	-36370.626	2742.785	994.993	264.686	-