

Table S2. Statistical assessment of the congruence between the Maximum Likelihood (ML) and Bayesian tree topologies.

<b>Tree of the concatenated nucleotides</b>	<b>-ln L<sup>a</sup></b>	<b>Diff -ln L<sup>b</sup></b>	<b>p-value</b>
ML tree topology	29555.84144		
Bayesian tree topology	29607.08844	51.24700	0.65