

Additional data file 10 - Comparison of performance of aLRT results and Bayesian posterior probabilities.

Comparison shows node support for kinesin phylogeny shown in Additional data file 2 inferred from Bayesian posterior probabilities (PP; 8 runs of 12,000,000 generations) or non-parametric Shimodaira-Hasegawa-like (SH) and parametric χ^2 -based (Chi2) approximate Likelihood Ratio Test.

