

FEL

Fixed Effects Likelihood

FEL analysis was performed on the alignment from `/home/datamonkey/datamonkey-js-server/production/app/fel/output/668ba39ffec0145ecad6d020`. Statistical significance is evaluated based on the asymptotic χ^2 distribution. This analysis **includes** site to site synonymous rate variation. .

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Suggested citation: Not So Different After All: A Comparison of Methods for Detecting Amino Acid Sites Under Selection (2005). *Mol Biol Evol* 22 (5): 1208-1222

p-value threshold [Update](#)

19

sequences in the alignment

619

codon sites in the alignment

1

partitions

33

median branches/partition used for testing

522

non-invariant sites tested

N/A

parametric bootstrap replicates

1Sites under diversifying positive selection at $p \leq 0.1$ **446**Sites under purifying selection at $p \leq 0.1$

Figure 1. Maximum likelihood estimates of synonymous (α) and non-synonymous rates (β) at each site. Estimates above 10 are censored at this value. p-values are also shown

