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URL: https://www.datamonkey.org/busted/5f45198134c53e423ae6751a

## Branch-site Unrestricted Statistical Test for Episodic Diversification results summary

INPUT DATA | 5f45198134c53e423ae6751a | 44 sequences | 561 sites

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BUSTED with synyonymous rate variation **found no evidence** (LRT, p-value =  $0.478 \ge .05$ ) of gene-wide episodic diversifying selection in the selected test branches of your phylogeny. Therefore, there is no evidence that any sites have experienced diversifying selection along the test branch(es).

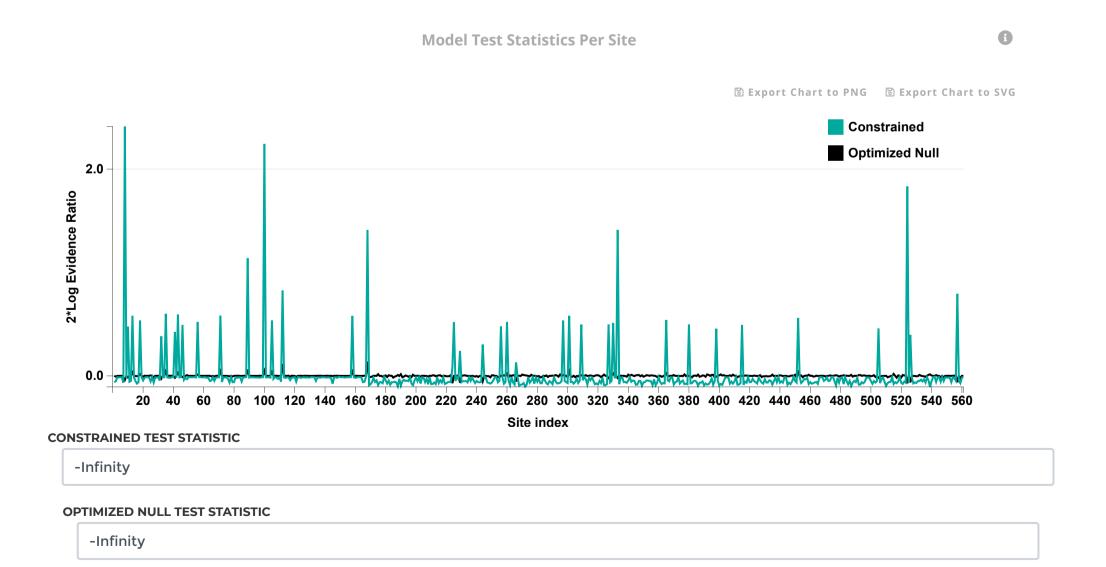
See **here** for more information about this method.

Please cite **PMID 25701167** if you use this result in a publication, presentation, or other scientific work.

Model log L AIC<sub>c</sub> CV(SRV) **Branch set** #. params  $\omega_1$  $\omega_2$  $\omega_3$ 0.01 (94.89%) 1.60 (1.53%) **Unconstrained model** -6150.5 109 12520.0 0.839 Test 0.02 (3.58%) dd 0.02 (6.99%) **Constrained model** -6150.6 108 12518.1 0.838 0.00 (90.04%) 1.00 (2.97%) Test dd

**Model fits** 

This table reports a statistical summary of the models fit to the data. Here, **Unconstrained model** refers to the BUSTED alternative model for selection, and **Constrained model** refers to the BUSTED null model for selection.



Showing entries 1 through 20 out of 561.

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Site index Unconstrained Constrained Optimized Null Constrained Optimized Null

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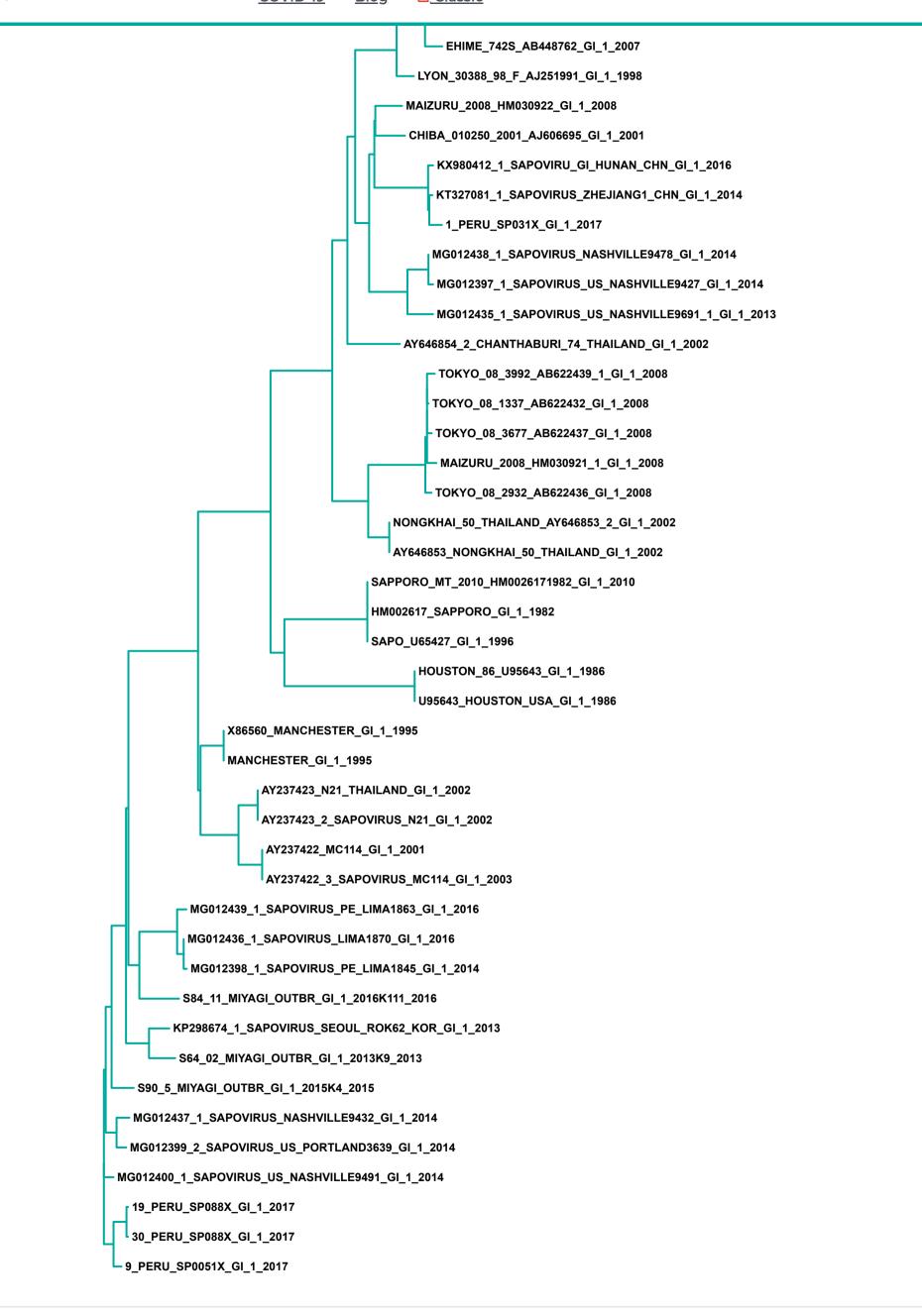
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2	-10.15	-10.13	-10.15	-0.05	-0.01
3	-5	-4.99	-5	-0.01	0
4	-4.94	-4.93	-4.94	-0.02	0
5	-5	-4.99	-5	-0.01	0
6	-4.92	-4.91	-4.92	-0.01	0
7	-13.39	-13.36	-13.38	-0.07	-0.02
8	-42.05	-43.25	-42.03	2.41	-0.04
9	-4.66	-4.65	-4.66	-0.02	0
10	-17.87	-18.1	-17.86	0.48	-0.02
11	-9.8	-9.77	-9.8	-0.06	0
12	-18.07	-18.04	-18.07	-0.05	0
13	-14.82	-15.11	-14.84	0.58	0.03
14	-4.89	-4.89	-4.89	-0.02	0
15	-10.35	-10.32	-10.35	-0.05	0
16	-15.8	-15.76	-15.79	-0.07	0
17	-4.24	-4.21	-4.24	-0.06	0
18	-17.16	-17.43	-17.17	0.54	0.02
19	-4.74	-4.74	-4.74	-0.01	0
20	-9.28	-9.26	-9.28	-0.04	-0.01

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Datamonkey is funded jointly by MIDAS and NIH award R01 GM093939





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