



Branch-site Unrestricted Statistical Test for Episodic Diversification results summary

INPUT DATA | [5f451a7a34c53e423ae67563](#) | 39 sequences | 570 sites

 Export ▾

BUSTED with synonymous rate variation **found evidence** (LRT, p-value = $0.000 \leq .05$) of gene-wide episodic diversifying selection in the selected test branches of your phylogeny. Therefore, there is evidence that at least one site on at least one test branch has experienced diversifying selection. 

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.

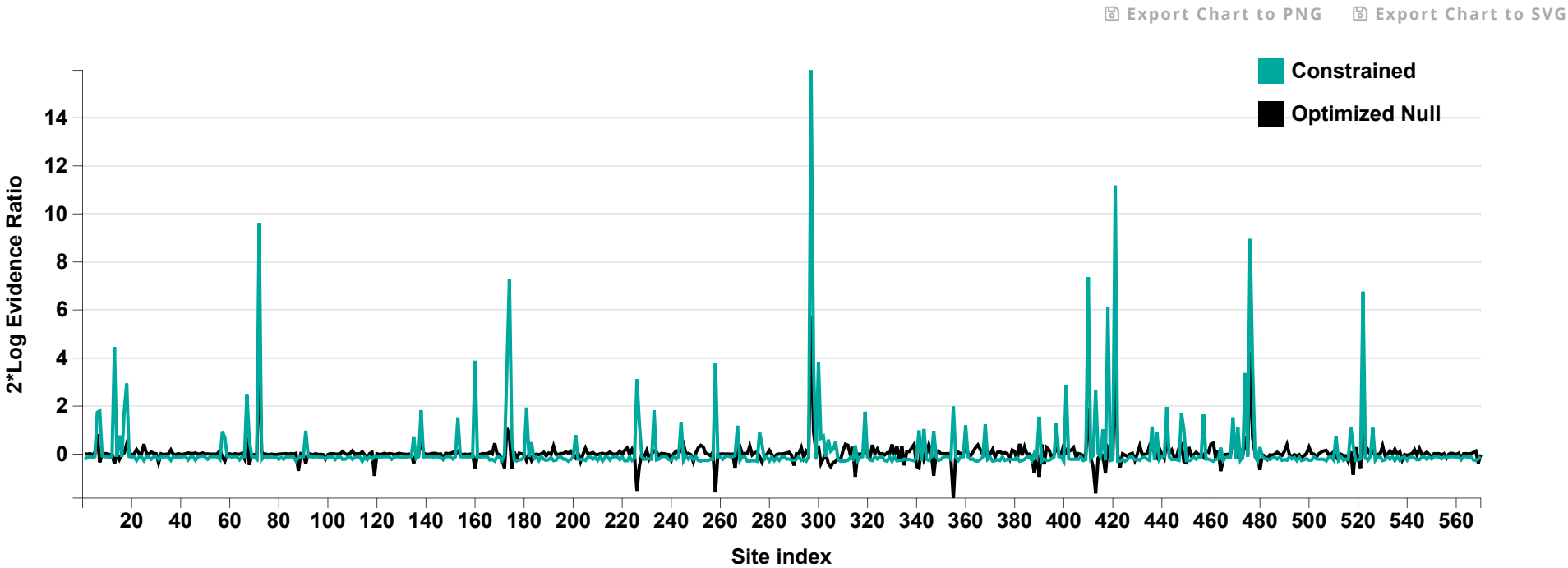
Model fits



Model	log L	#. params	AIC _c	CV(SRV)	Branch set	ω ₁	ω ₂	ω ₃	
Unconstrained model	-5510.3	99	11219.4	0.737	Test	0.03 (1.18%)	0.06 (98.12%)	18.81 (0.70%)	Link
Constrained model	-5520.0	98	11236.9	0.754	Test	0.00 (13.20%)	0.00 (75.35%)	1.00 (11.45%)	Link

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.

Model Test Statistics Per Site



CONSTRAINED TEST STATISTIC

-Infinity

OPTIMIZED NULL TEST STATISTIC

-Infinity

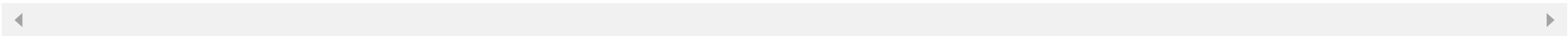
Showing entries 1 through 20 out of 570.

 Export Table to CSV

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Site indexUnconstrainedConstrainedOptimized NullConstrainedOptimized Null

2	-4.75	-4.69	-4.75	-0.12	0
3	-4.98	-4.93	-4.99	-0.1	0.02
4	-4.59	-4.52	-4.58	-0.13	-0.02
5	-4.98	-4.93	-4.99	-0.1	0.02
6	-16.58	-17.44	-16.99	1.74	0.83
7	-23.05	-23.95	-22.88	1.8	-0.34
8	-5.45	-5.41	-5.45	-0.07	0.01
9	-5.11	-5.06	-5.12	-0.11	0.02
10	-4.83	-4.77	-4.84	-0.11	0.02
11	-4.59	-4.52	-4.58	-0.13	-0.02
12	-5.03	-4.98	-5.03	-0.11	-0.01
13	-34.27	-36.5	-34.07	4.46	-0.41
14	-4.72	-4.66	-4.72	-0.11	0
15	-19.03	-19.42	-18.92	0.78	-0.22
16	-4.78	-4.72	-4.77	-0.12	-0.02
17	-14.79	-15.7	-14.82	1.82	0.06
18	-21.92	-23.39	-22.13	2.94	0.42
19	-5.17	-5.12	-5.17	-0.1	0
20	-4.32	-4.26	-4.31	-0.12	-0.02



Fitted tree

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Options ▾

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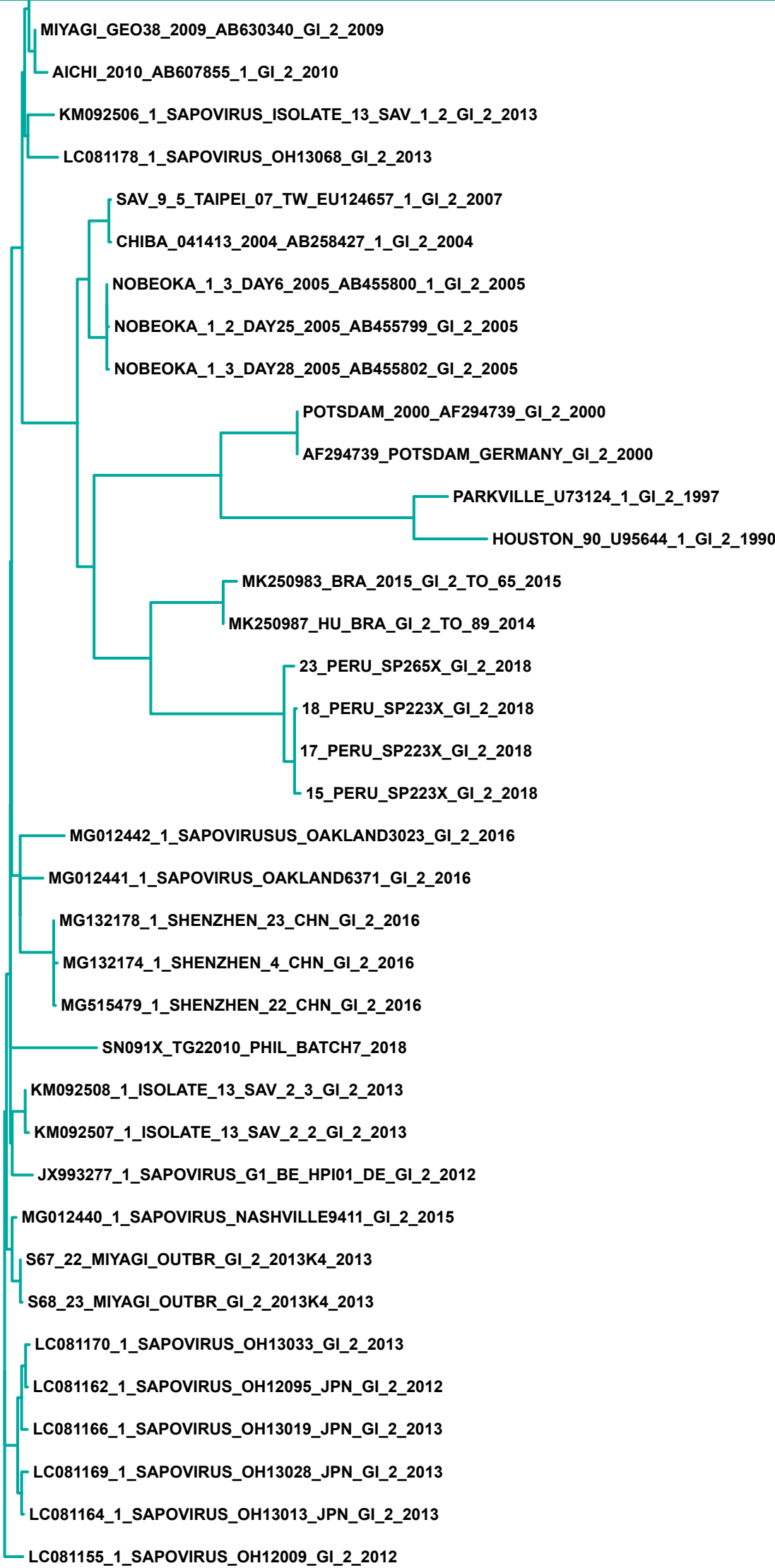
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Steven Weaver
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I've started working on a comprehensive benchmarking notebook for HyPhy and started with commonly used methods. If there are any specific benchmarks you'd like to see, let me know and I'll include if reasonable.

[observablehq.com/@stevenweaver/...](https://observablehq.com/@stevenweaver/)

HyPhy Benchmarks and Profiling
An Observable notebook by Steven Wea...
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