

Multi-Hit
results summary

INPUT DATA | WestNileVirus_NS3.fas | 19 sequences | 619 sites

Export

Likelihood Test Results

3H vs 1H

LRT 7.406

p-value 0.060

3H vs. 2H

LRT 0.006

p-value 0.997

3H vs 3HSI

LRT -0.000

p-value 1.000

2H vs 1H

LRT 7.400

p-value 0.007

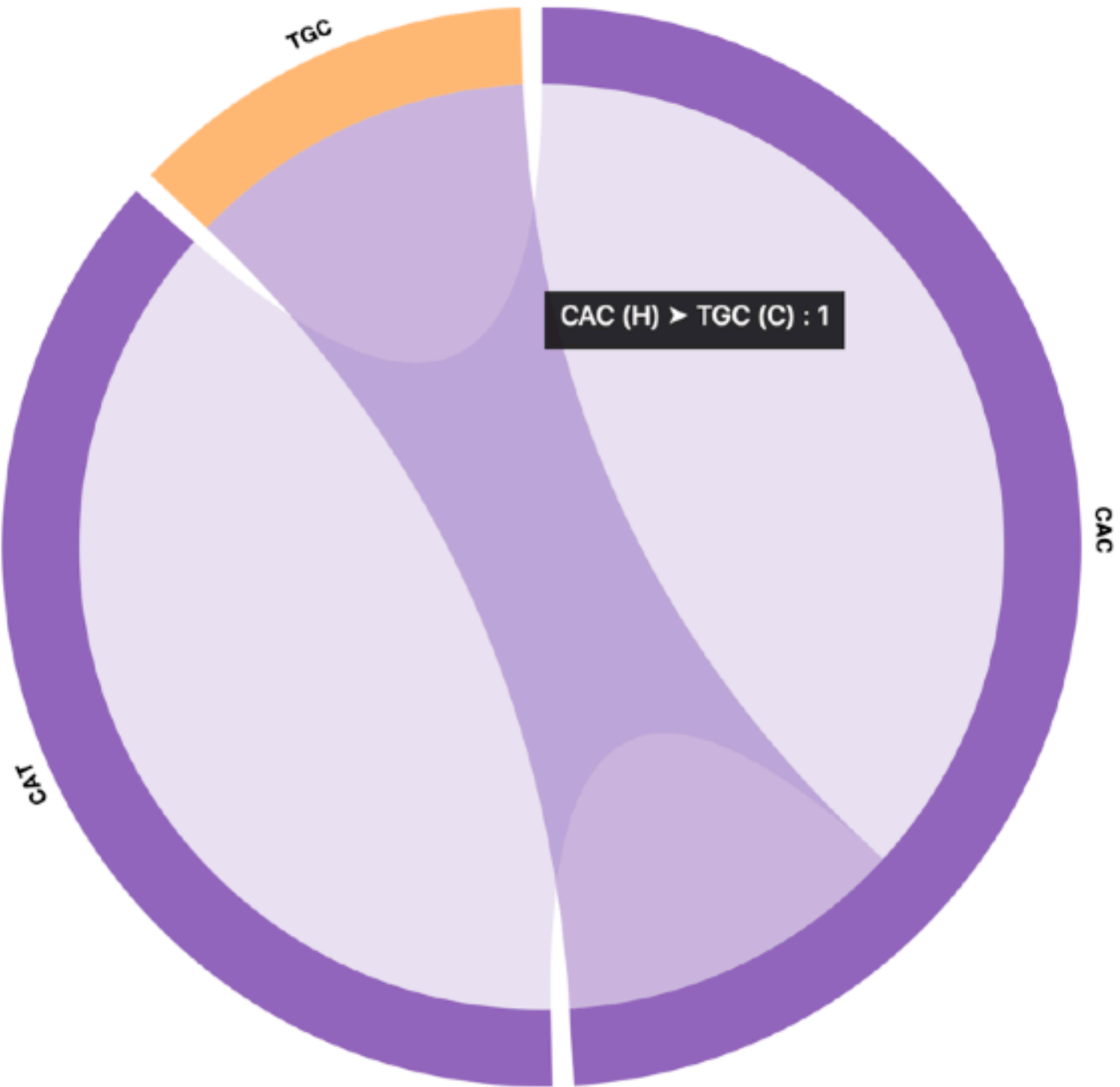
3HSI vs 2H

LRT 0.006

p-value 0.936

See [here](#) for more information about this method.

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1 individual site which showed sufficiently strong preference for multiple-hit models

Site	ER (2 vs 1)	ER (3 vs 2)	ER (3-island vs 2)	ER (3-island vs 3)	Substitutions
87	54.0092	1.0065	1.0020	1.0045	CAC->CAT(3)TGC(1)

hyphy fmm --alignment WestNileVirus_NS3.fas --triple-islands Yes

