

# FEL

Fixed Effects Likelihood

Choose File No file chosen

Genetic Code<sup>?</sup> Universal code ▾

Synonymous rate variation (recommended)? Yes ▾

## NOTIFY WHEN COMPLETED?

Email Address

Advanced Options

Run Analysis ▶

Datamonkey is funded jointly by **MIDAS** and **NIH award R01 GM093939**

# FEL

Fixed Effects Likelihood

Choose File WestNileVirus\_NS3.fna

Genetic Code<sup>?</sup> Universal code ▾

Synonymous rate variation (recommended)? Yes ▾

## NOTIFY WHEN COMPLETED?

Email Address

### [Advanced Options](#)

[Advanced setting, will result in MUCH SLOWER run time] Perform parametric bootstrap resampling to derive site-level null LRT distributions up to this many replicates per site. Recommended use for small to medium (<30 sequences) datasets.

Resample? 50

Compute profile likelihood confidence intervals for each variable site

Yes? ☐

Run Analysis ▶