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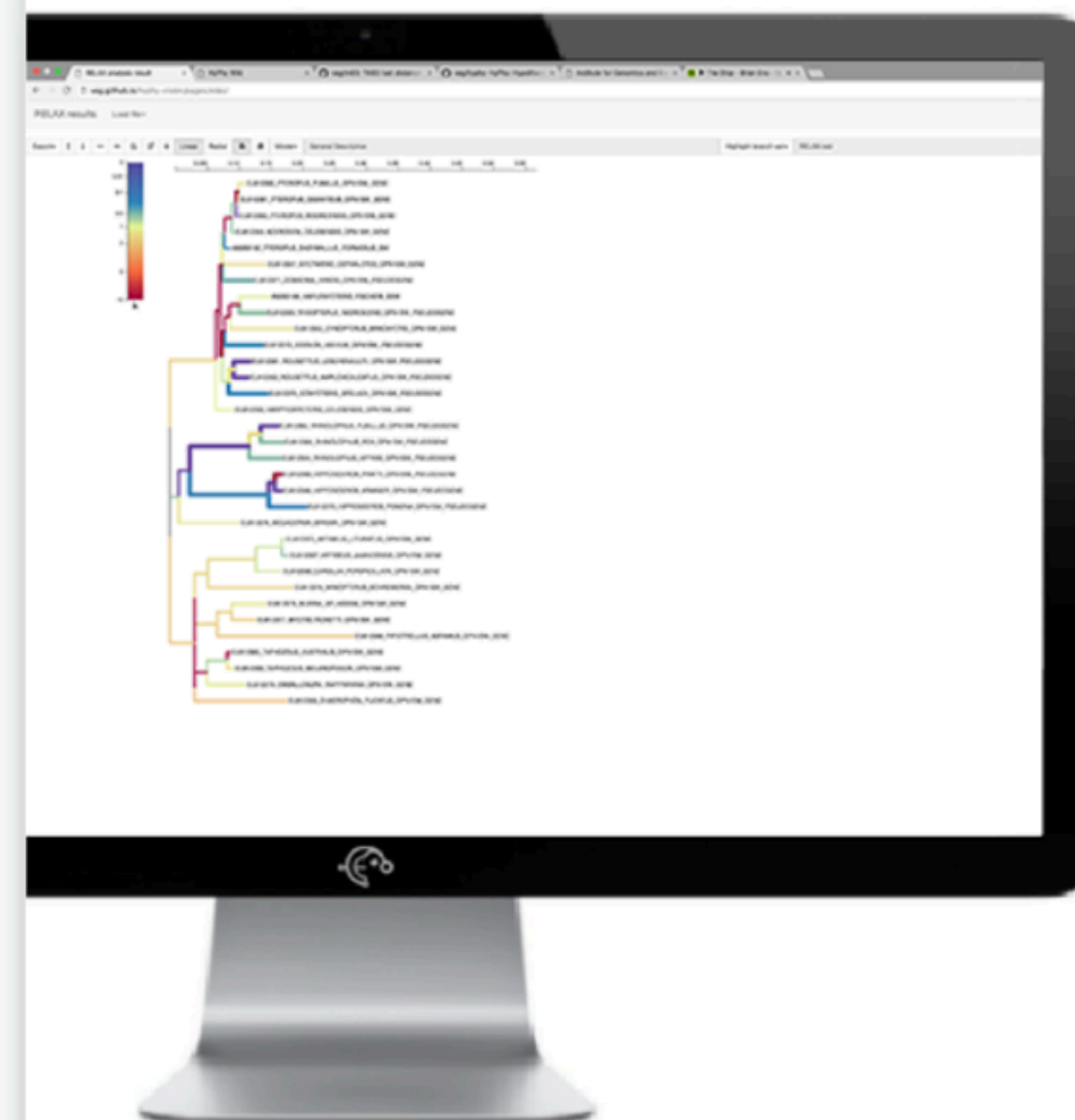
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Hypothesis Testing using Phylogenies

An open-source software package

for comparative sequence analysis
using stochastic evolutionary models

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Datamonkey

A Collection of State of the Art Statistical Models and Bioinformatics Tools

What evolutionary process would you like to detect?

Selection

Recombination

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