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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

## Would you like to detect selection across branches, individual sites, or an entire gene?

Branches Sites Gene

## Do you want to detect episodic or pervasive selection?

Episodic Pervasive

## Datamonkey recommends that you use...



MEME (Mixed EffectsModel of Evolution) employs a mixed-effects maximum likelihood approach to test the hypothesis that individual sites have been subject to episodic positive or diversifying selection. In other words, MEME aims to detect sites evolving under positive selection under a proportion of branches.

For more information, please see the <u>summary on hyphy.org</u> or see <u>Murrell, B et al. "Detecting individual sites subject to episodic diversifying selection." PLoS Genetics 8, e1002764 (2012).</u>

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https://datamonkey.org