phylogenies taxonomy unified tree of life teaching automated and outreach analyses community other reasearch composition tracing trait applications evolution

Emily Jane McTavish

Call me: 'Emily Jane' (she/her)

How can we take advantage of evolutionary information in large data sets, in a way that is:

- biologically informed
- statistically rigorous
- computationally efficient?



Twitter: @snacktavish GitHub: snacktavish

Website: https://mctavishlab.github.io/





