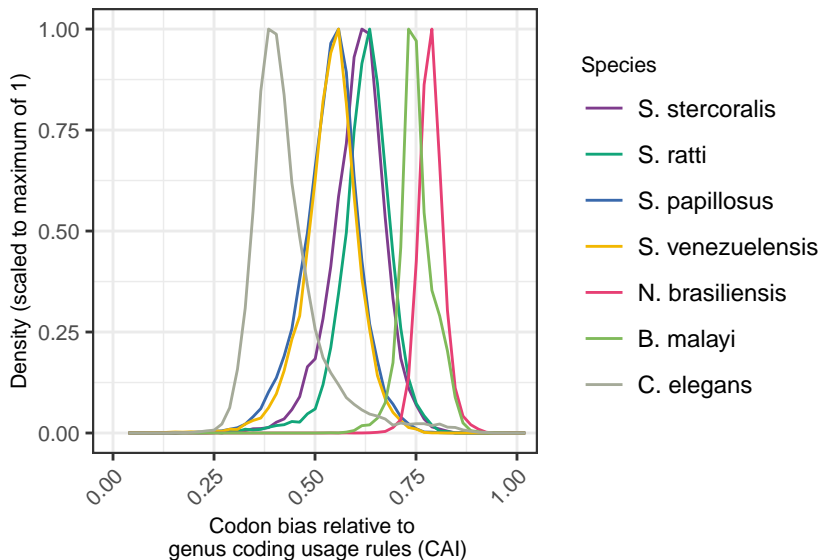


## Species-wide codon adaptiveness

Data includes all predicted coding sequences



Strongyloides species values relative to *S. ratti* usage rules

*C. elegans* values relative to *C. elegans* usage rules.

*N. brasiliensis* values relative to *N. brasiliensis* usage rules.

*B. malayi* values relative to *B. malayi* usage rules.