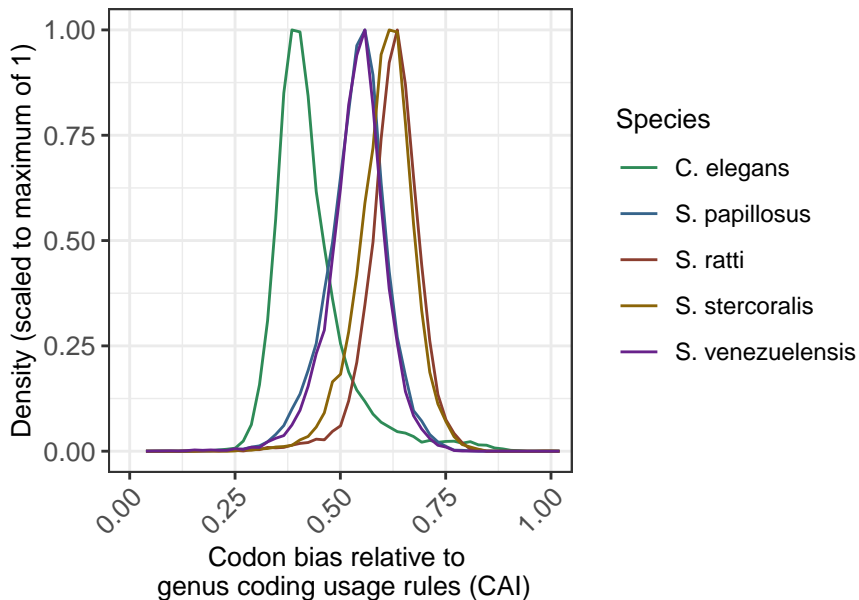


# Species-wide codon adaptiveness

Data includes all predicted coding sequences



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