

(a) Arctiina

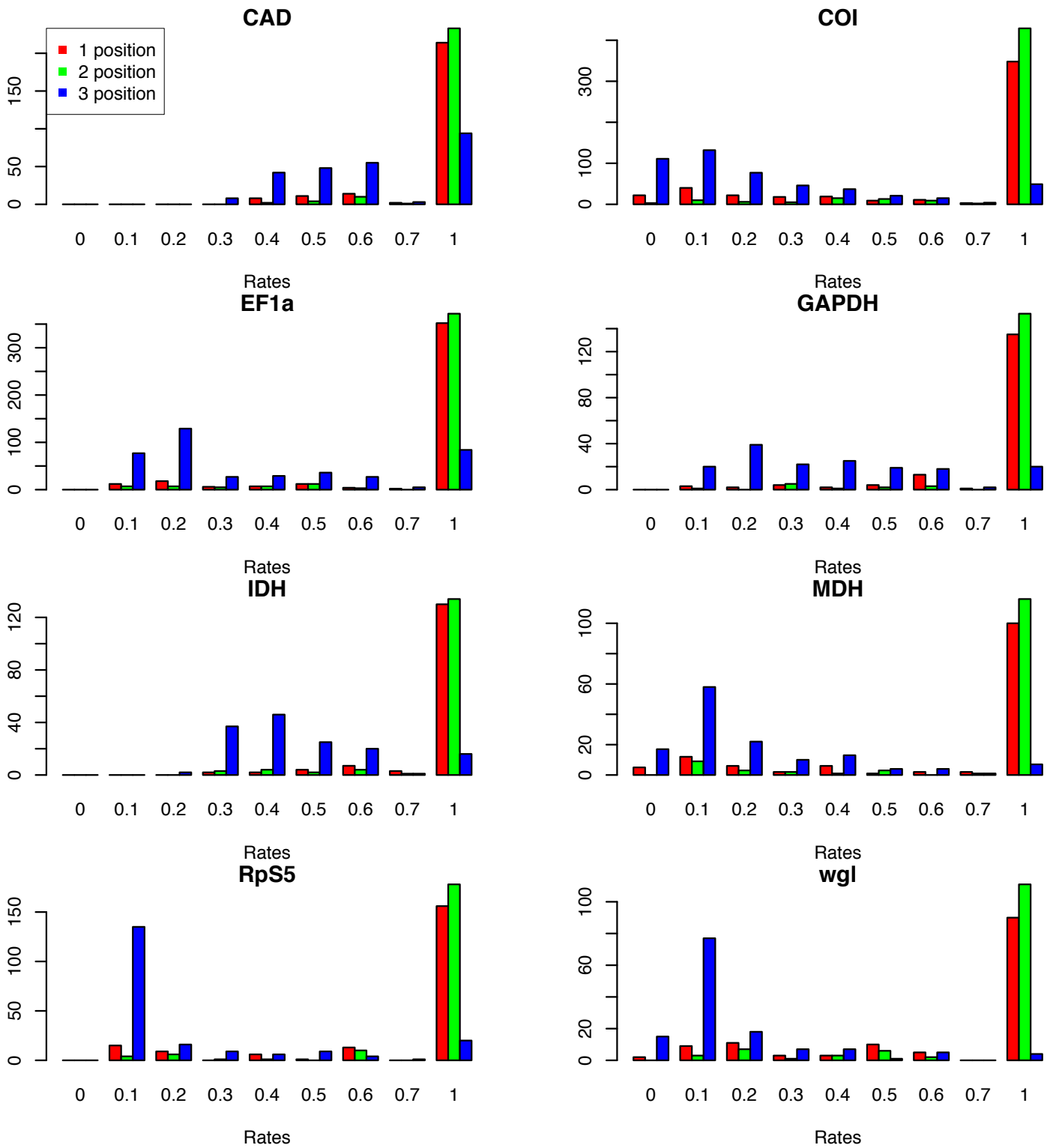
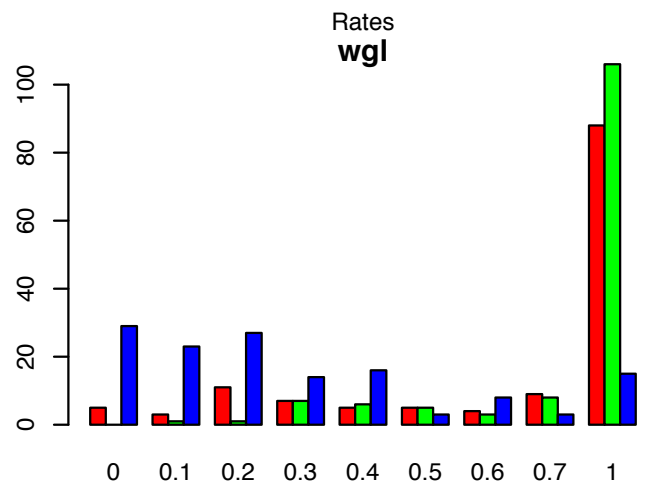
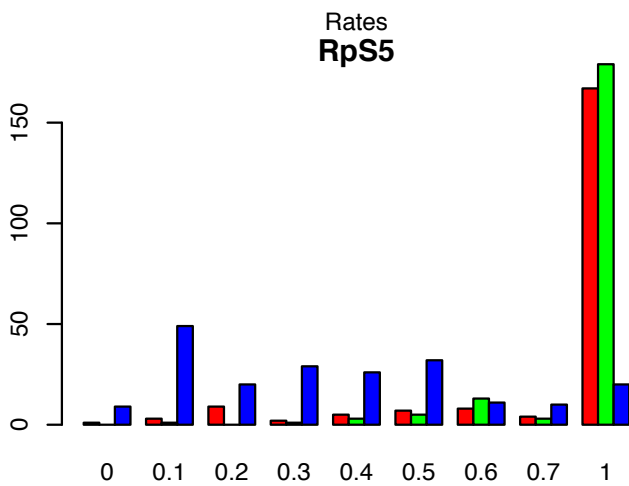
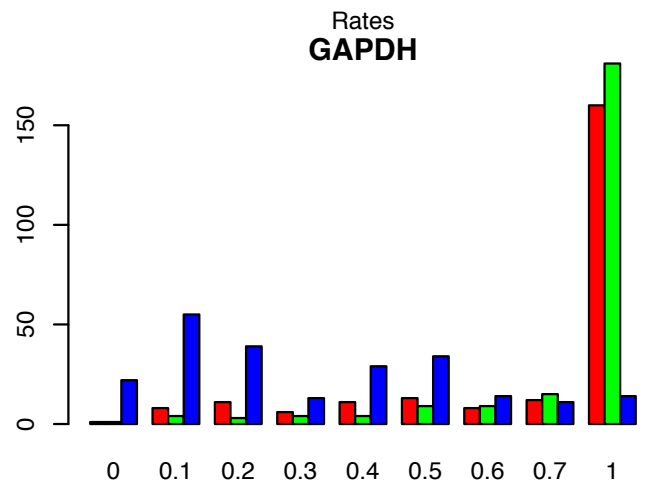
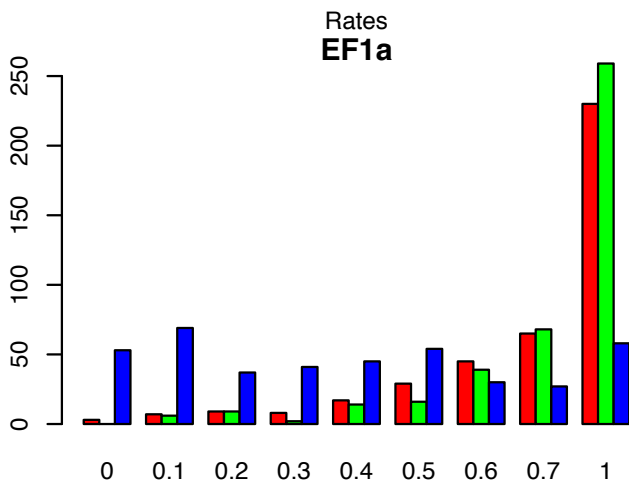
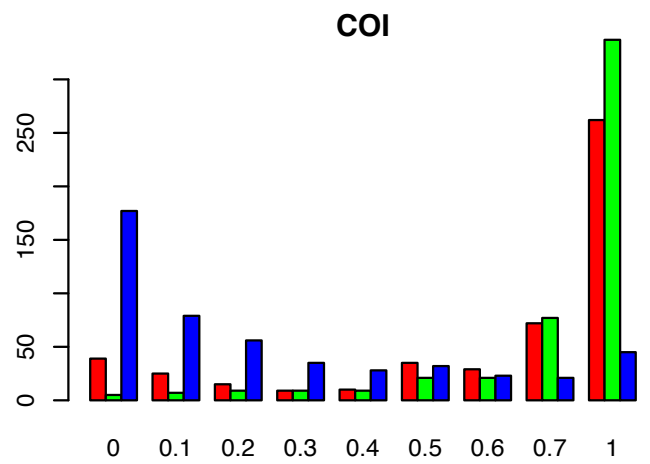
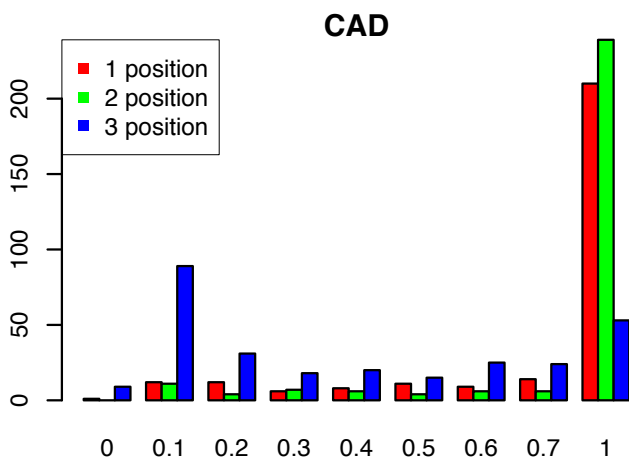


Figure S1. Plots showing the assigned TIGER rates to codon positions of each of the genes in datasets Arctiina (a), *Calisto* (b), Choreutidae (c), Coenonymphina (d), Geometridae (e), *Morpho* (f), and Pieridae (g).

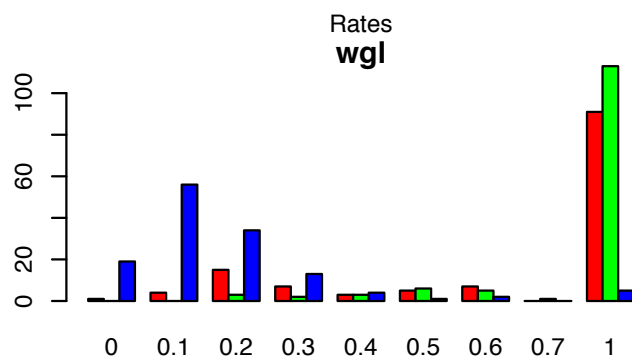
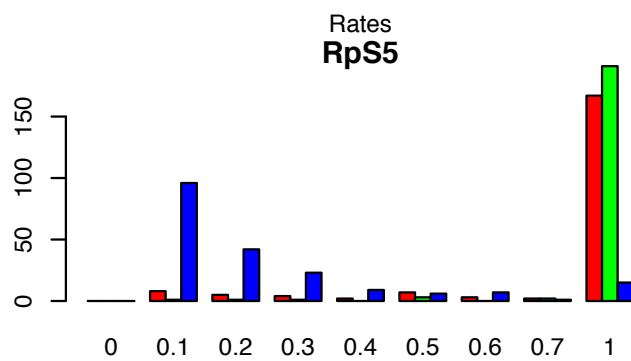
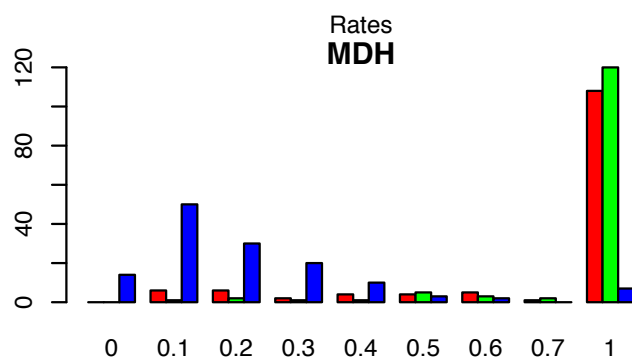
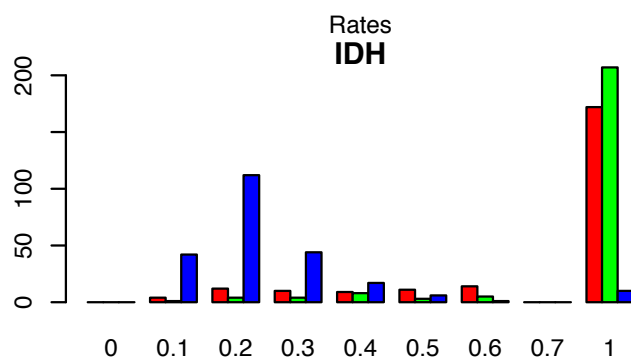
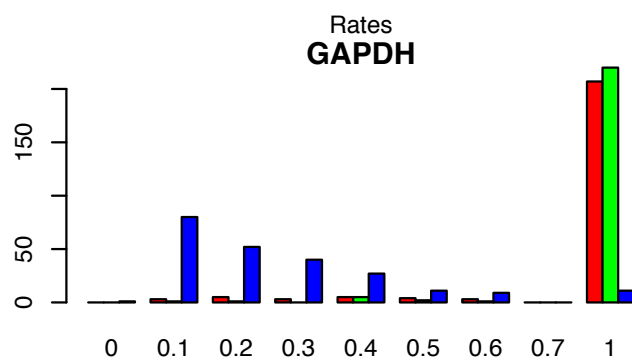
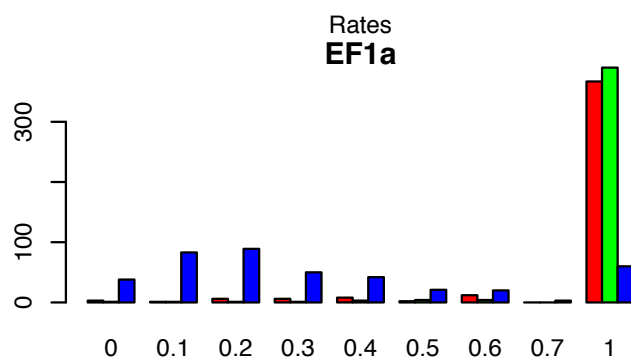
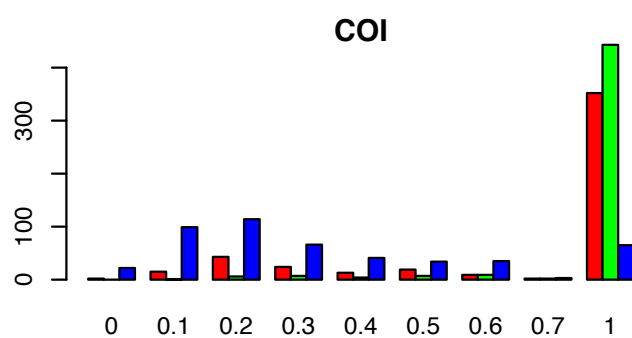
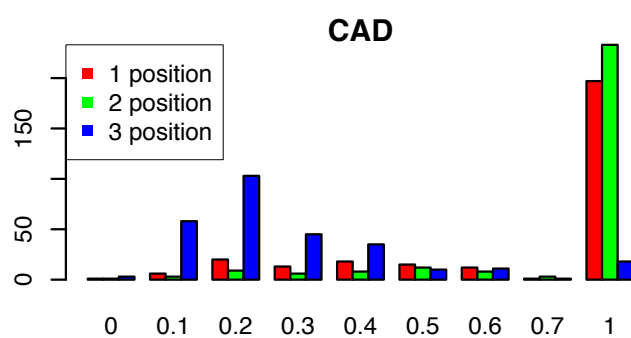
(b) Calisto



Rates

Rates

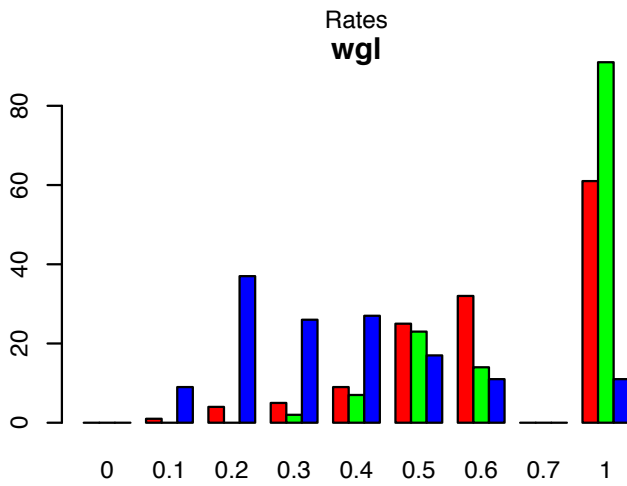
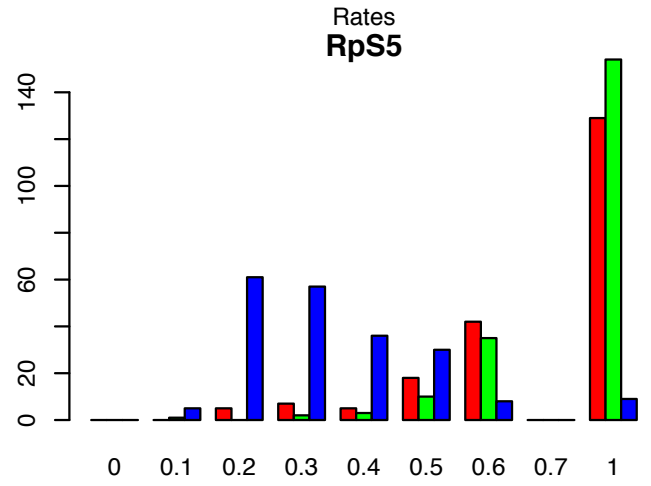
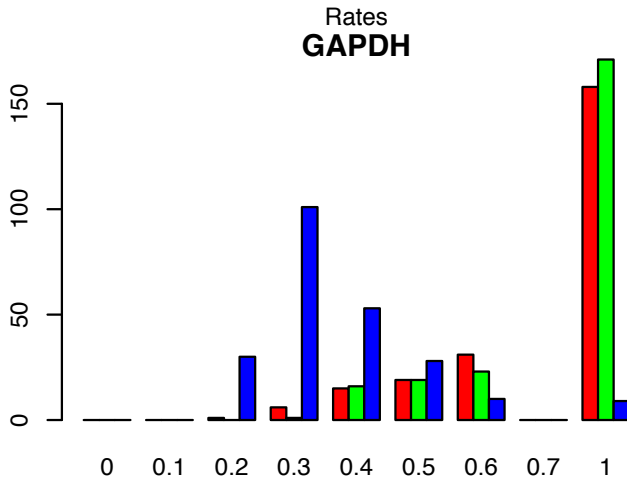
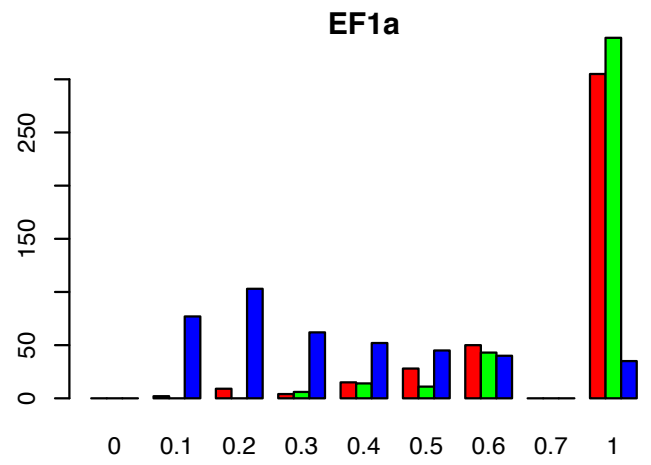
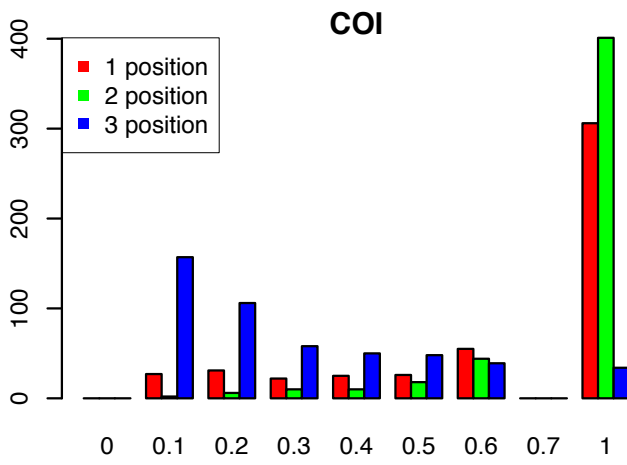
(c) Choreutidae



Rates

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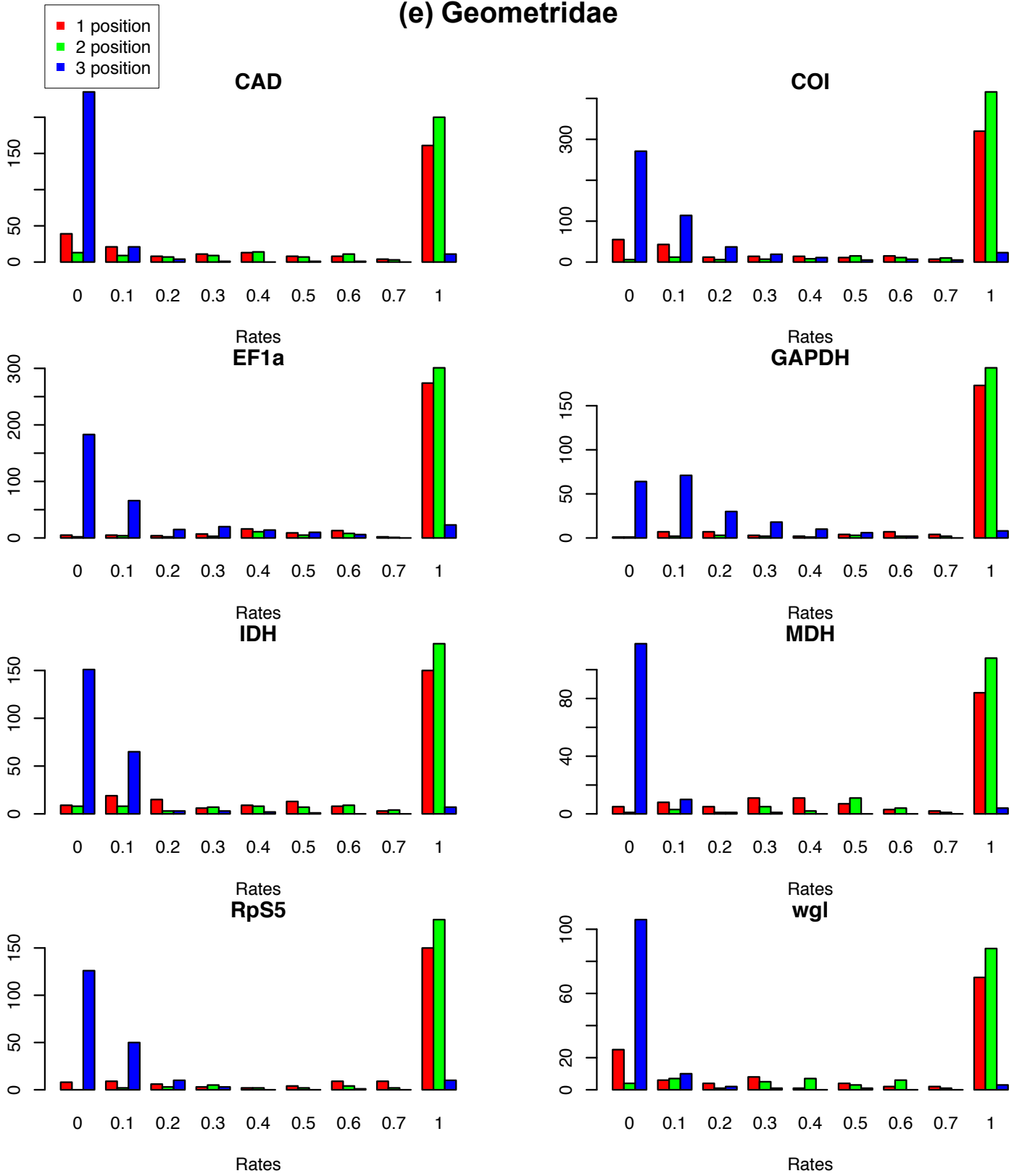
(d) Coenonymphina



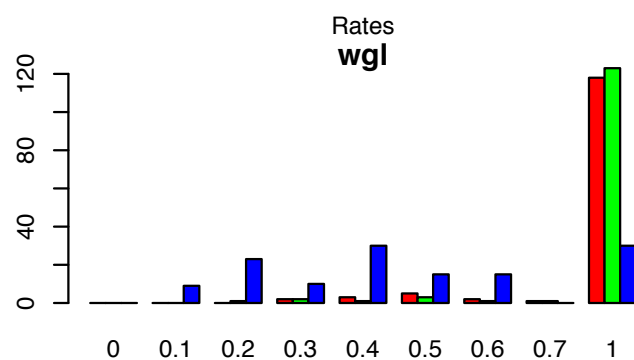
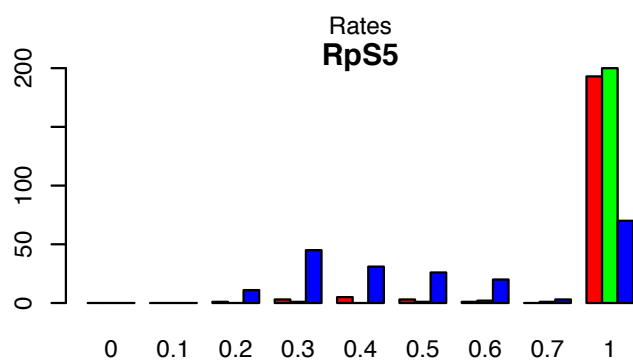
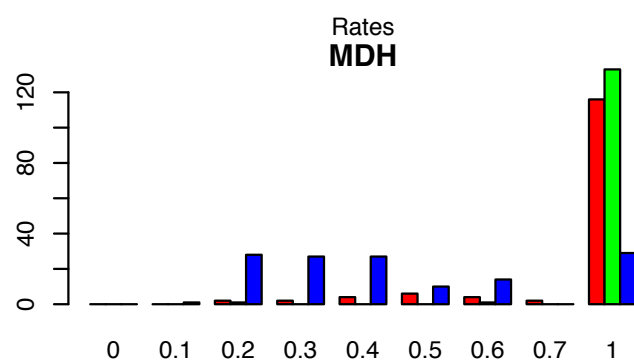
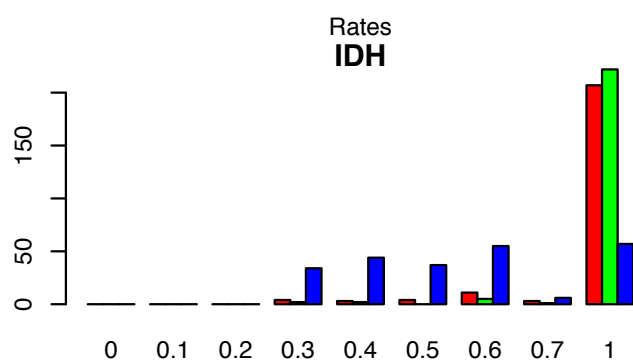
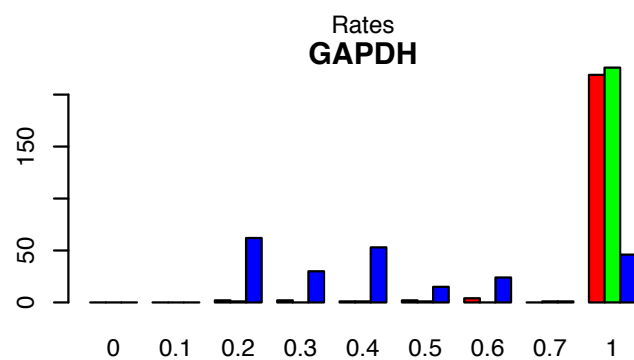
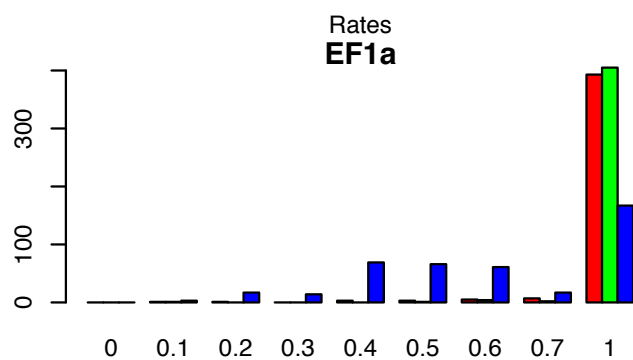
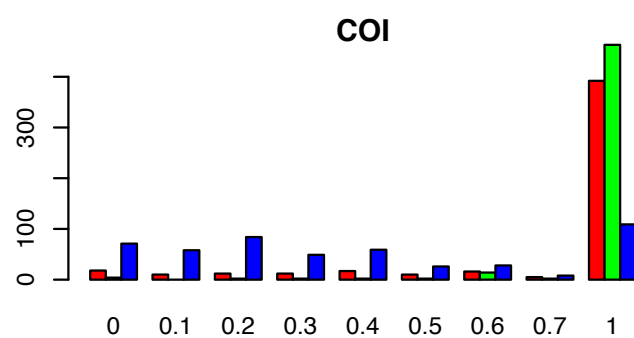
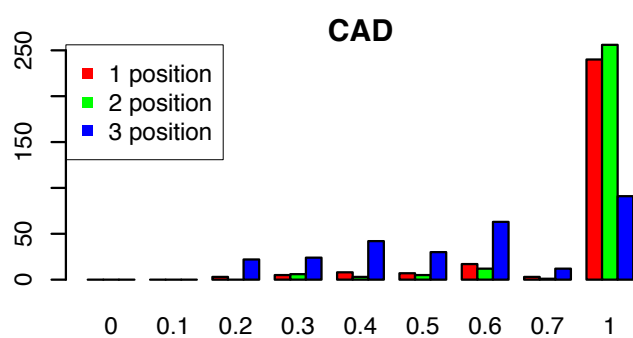
Rates

Rates

(e) Geometridae



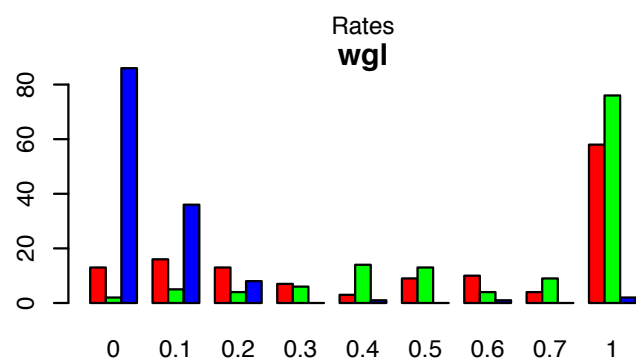
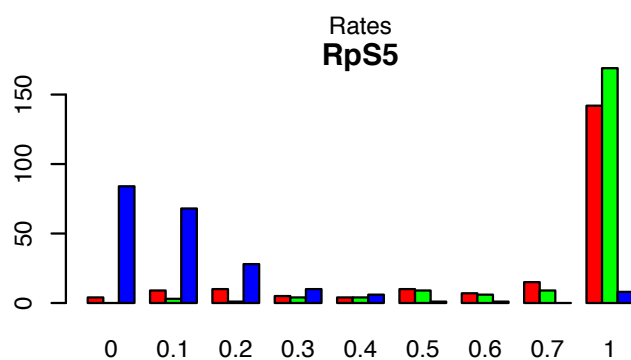
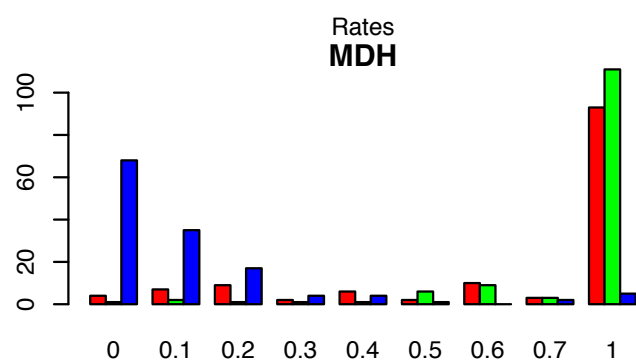
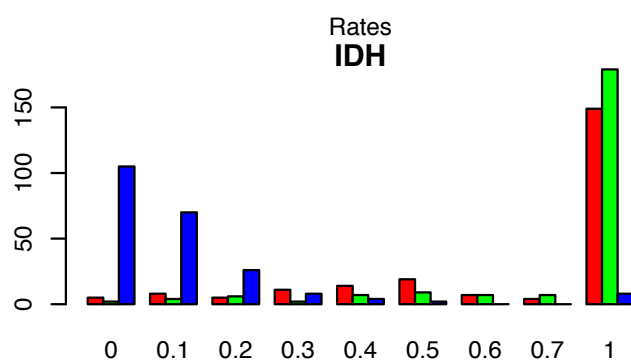
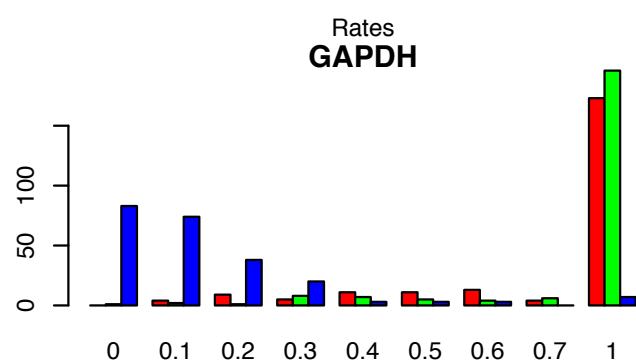
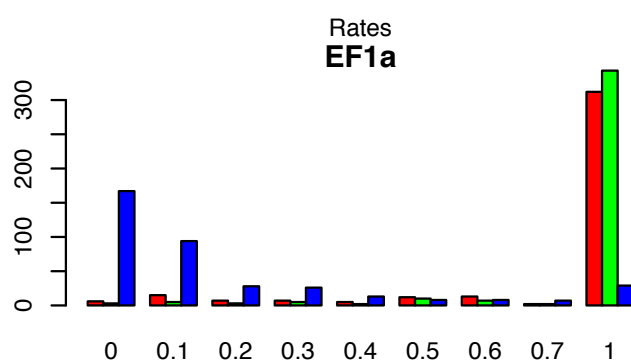
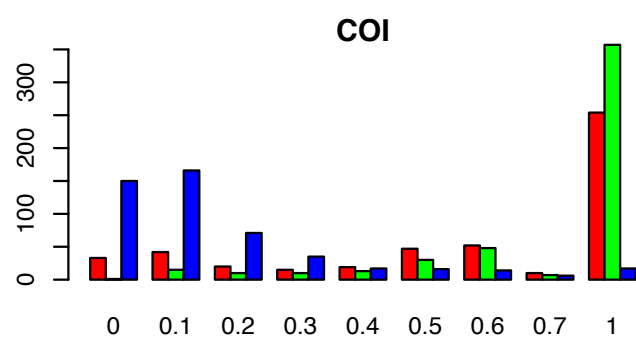
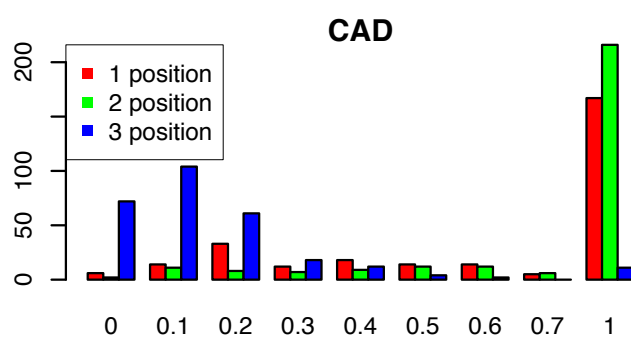
(f) *Morpho*



Rates

Rates

(g) Pieridae



Rates

Rates