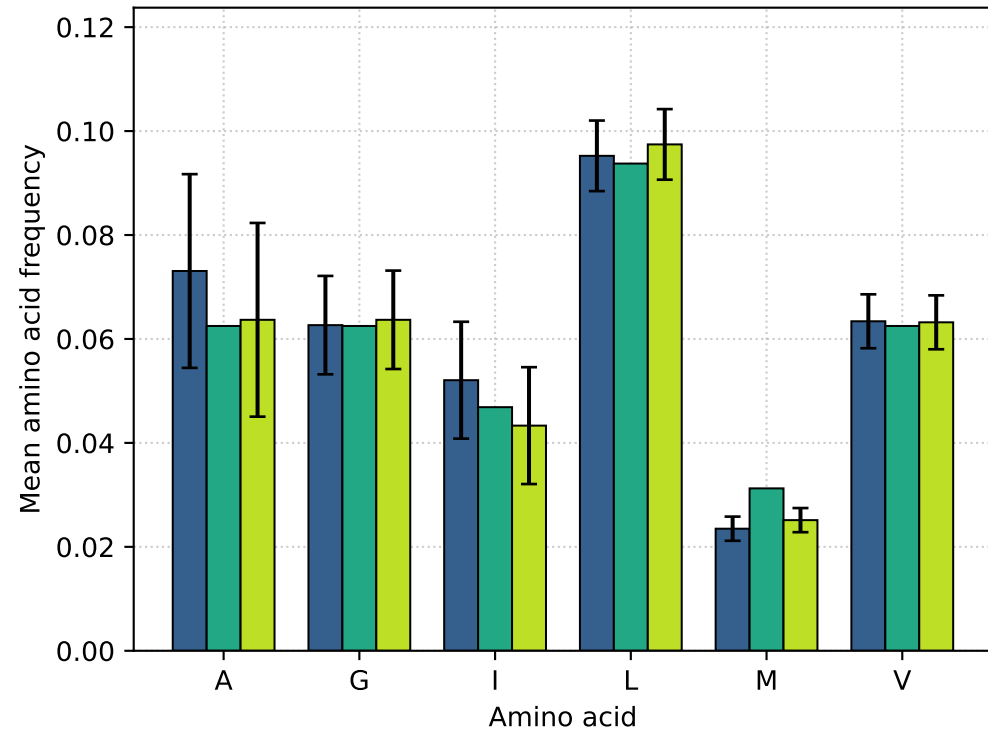
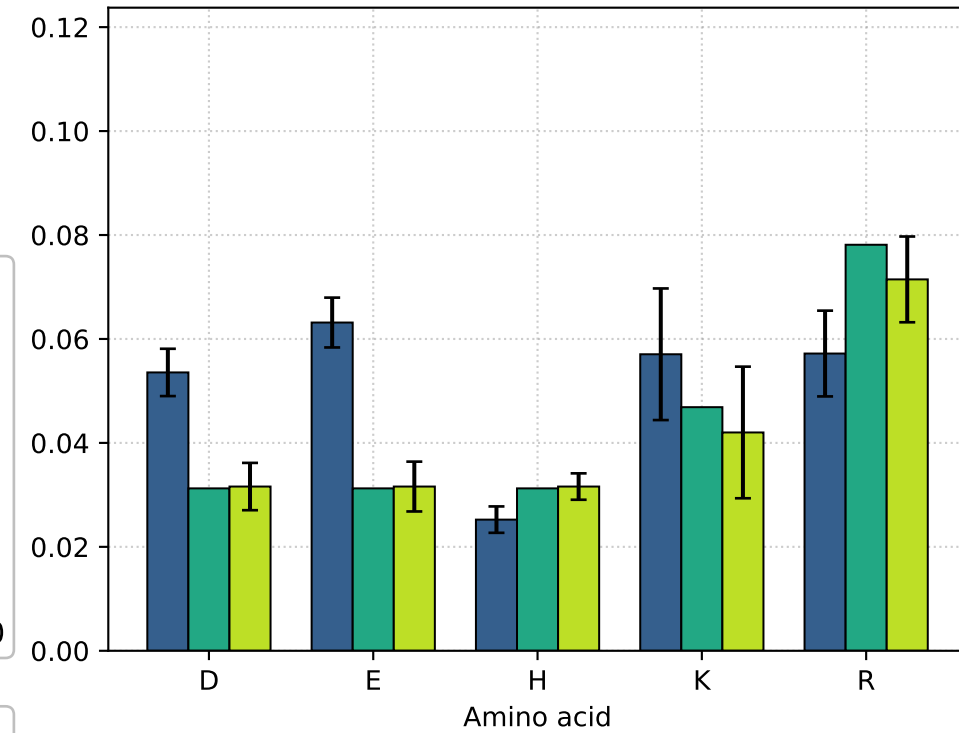


Eukaryota - Mean proteome amino acid frequency for genetic code: Rhabdopleuridae mitochondrial

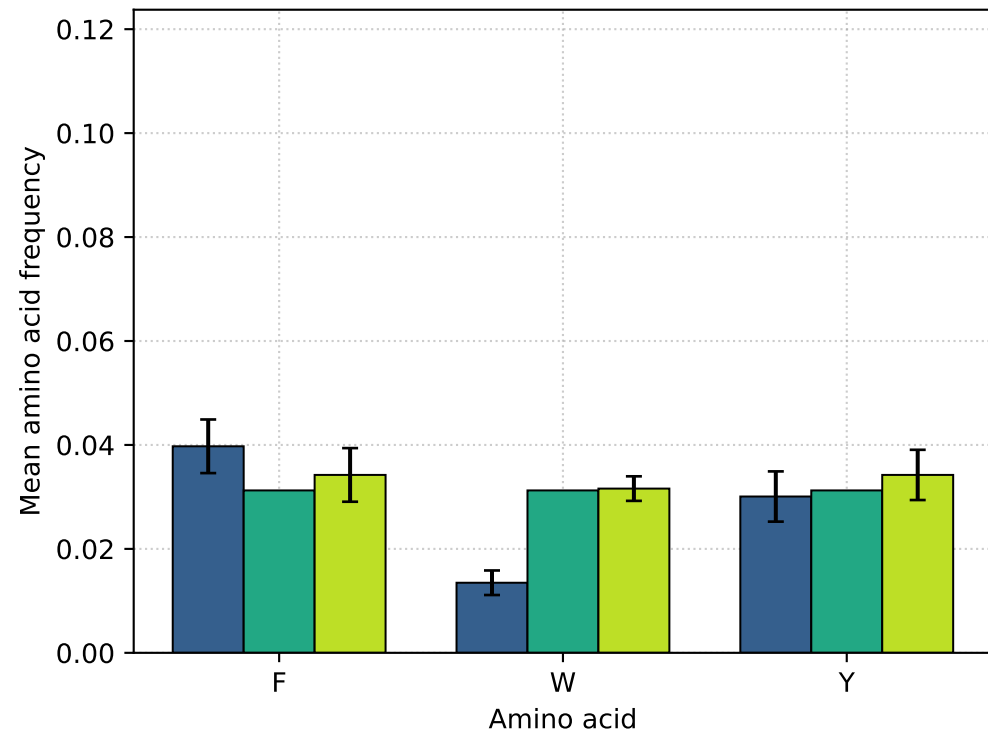
Aliphatic amino acids



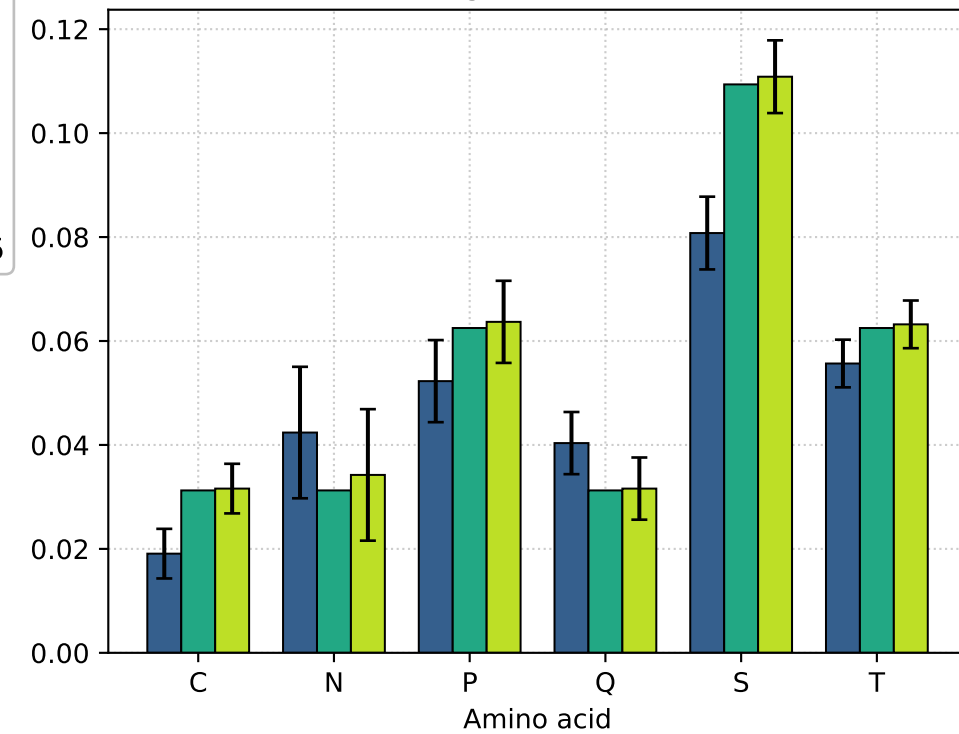
Charged amino acids



Aromatic amino acids



Uncharged amino acids



Observed
Codon
Codon+GC

Codon correlation:

- Mean Pearson:
- Coefficient: 0.75076
- p-value: 1.13853e-02

- Mean Spearman:
- Coefficient: 0.74317
- p-value: 1.27501e-02

- Mean log-MSE: 0.30590

Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.78968
- p-value: 2.13961e-04

- Mean Spearman:
- Coefficient: 0.75862
- p-value: 4.47813e-04

- Mean log-MSE: 0.29745