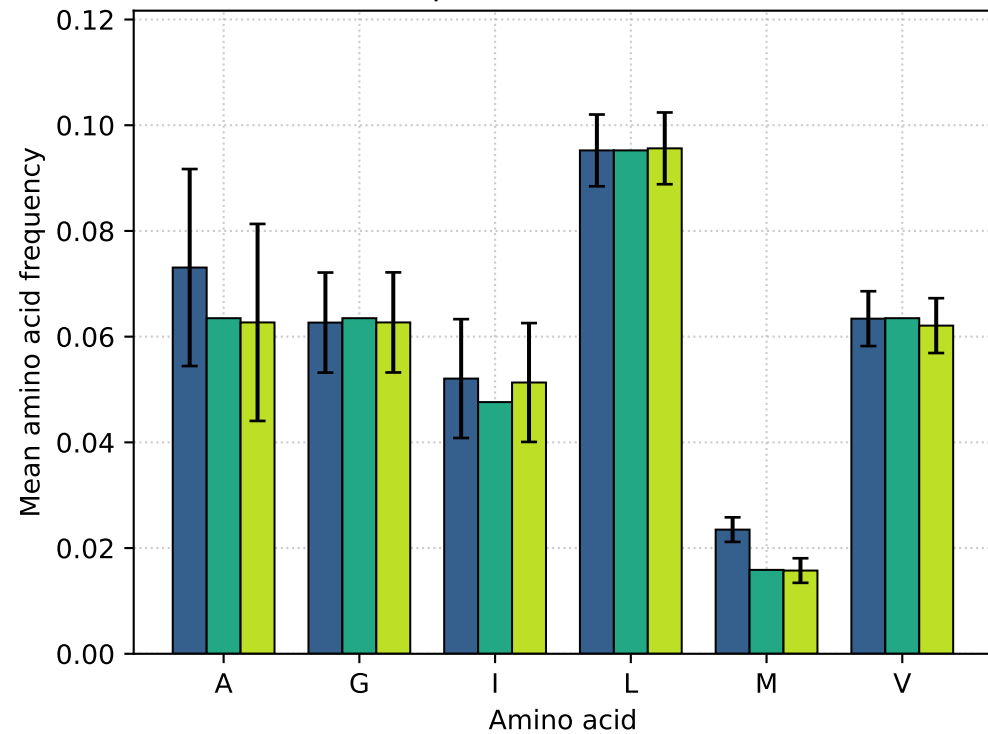
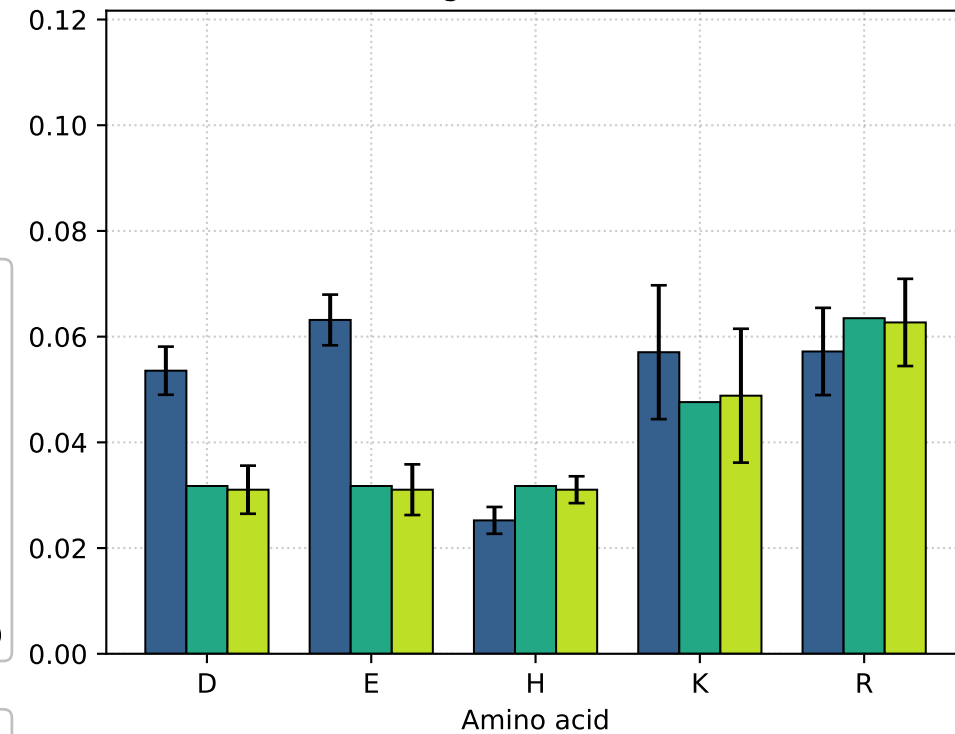


Eukaryota - Mean proteome amino acid frequency for genetic code: Cephalodiscidae mitochondrial uaa-tyr

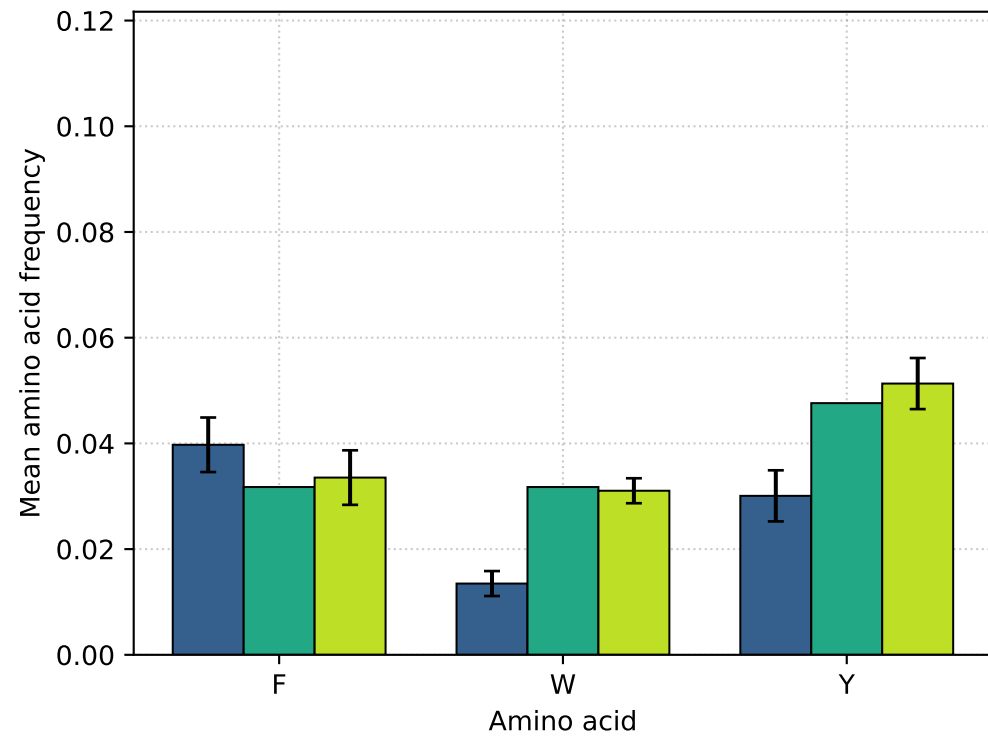
Aliphatic amino acids



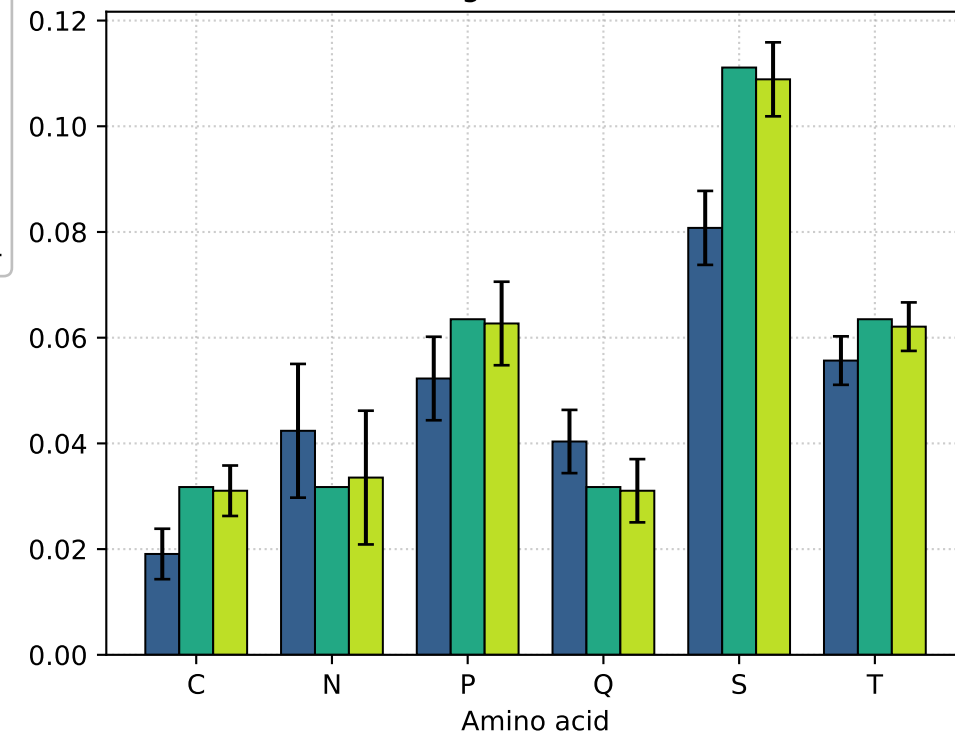
Charged amino acids



Aromatic amino acids



Uncharged amino acids



Codon correlation:

- Mean Pearson:
- Coefficient: 0.75555
- p-value: 8.31128e-03
- Mean Spearman:
- Coefficient: 0.73087
- p-value: 1.08982e-02
- Mean log-MSE: 0.32599

Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.78037
- p-value: 2.84351e-04
- Mean Spearman:
- Coefficient: 0.73588
- p-value: 8.07044e-04
- Mean log-MSE: 0.32074