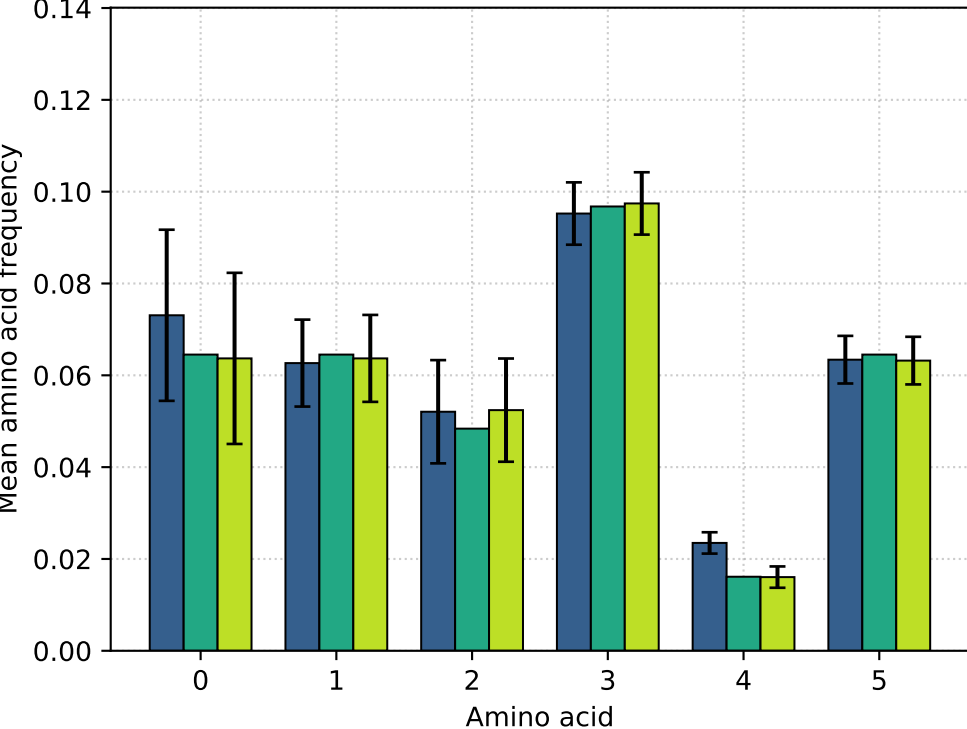
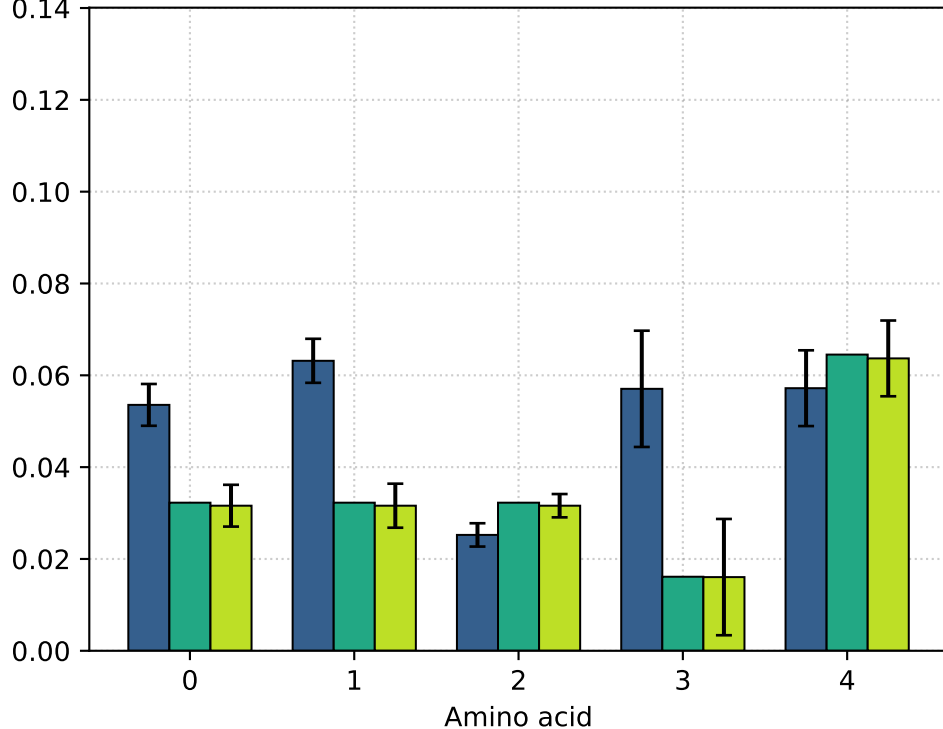


Eukaryota - Mean proteome amino acid frequency for genetic code: Echinoderm-flatworm mitochondrial

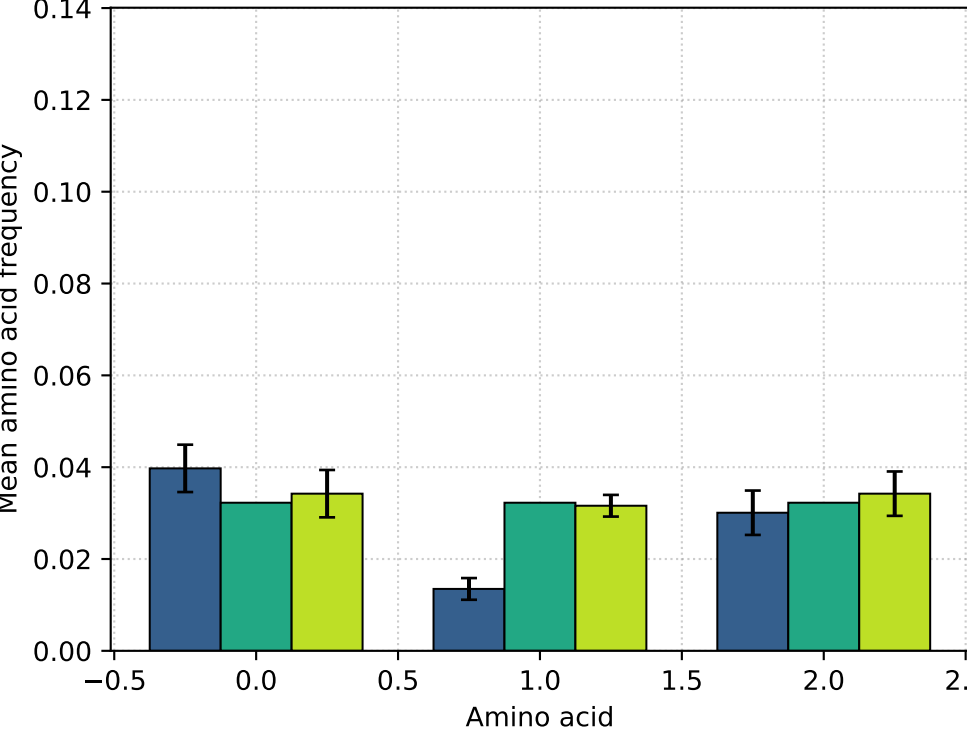
Aliphatic amino acids



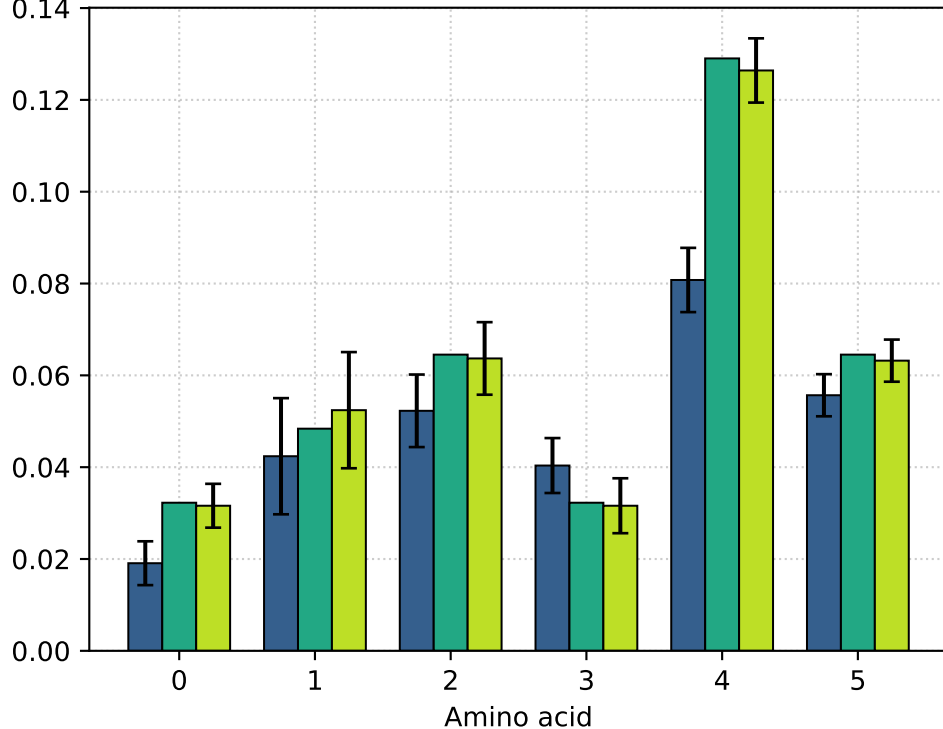
Charged amino acids



Aromatic amino acids



Uncharged amino acids



Legend:

- Observed
- Codon
- Codon+GC

Codon correlation:

- Mean Pearson:
- Coefficient: 0.69925
- p-value: 1.36370e-02
- Mean Spearman:
- Coefficient: 0.66282
- p-value: 2.07163e-02
- Mean log-MSE: 0.46971

Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.73167
- p-value: 9.28982e-04
- Mean Spearman:
- Coefficient: 0.66326
- p-value: 6.17076e-03
- Mean log-MSE: 0.46613