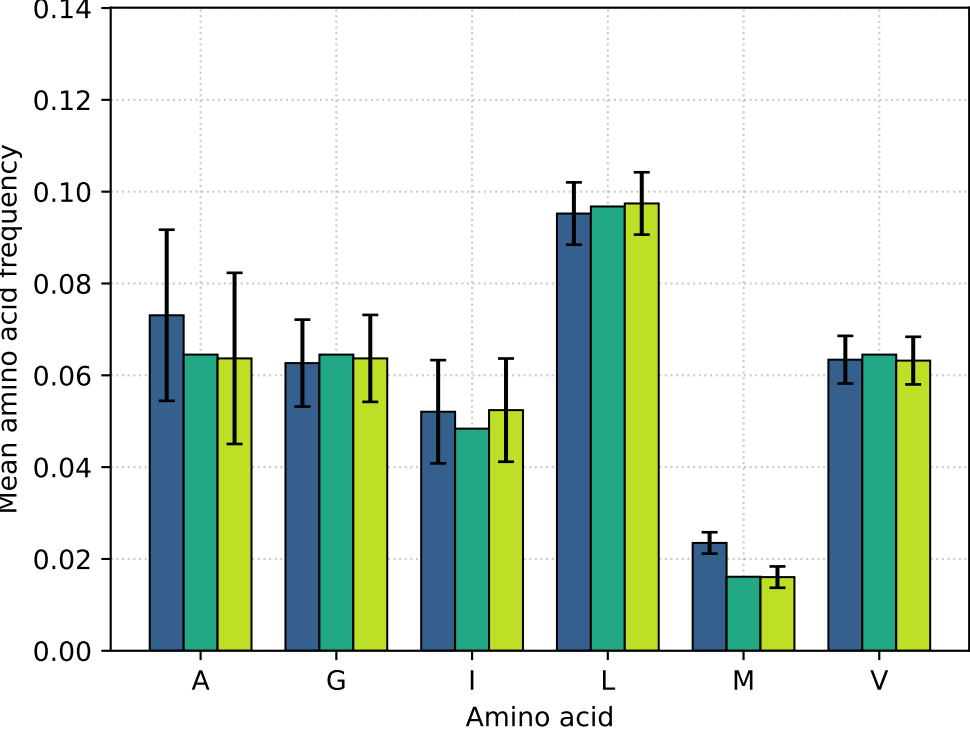
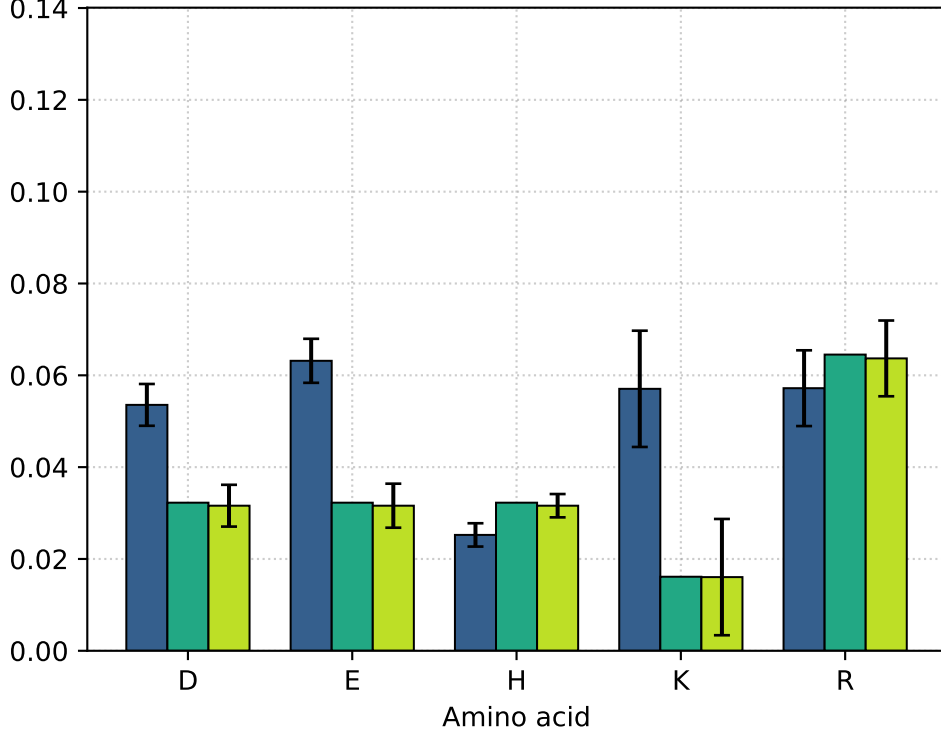


Eukaryota - Mean proteome amino acid frequency for genetic code: Echinoderm-flatwormmitochondrial

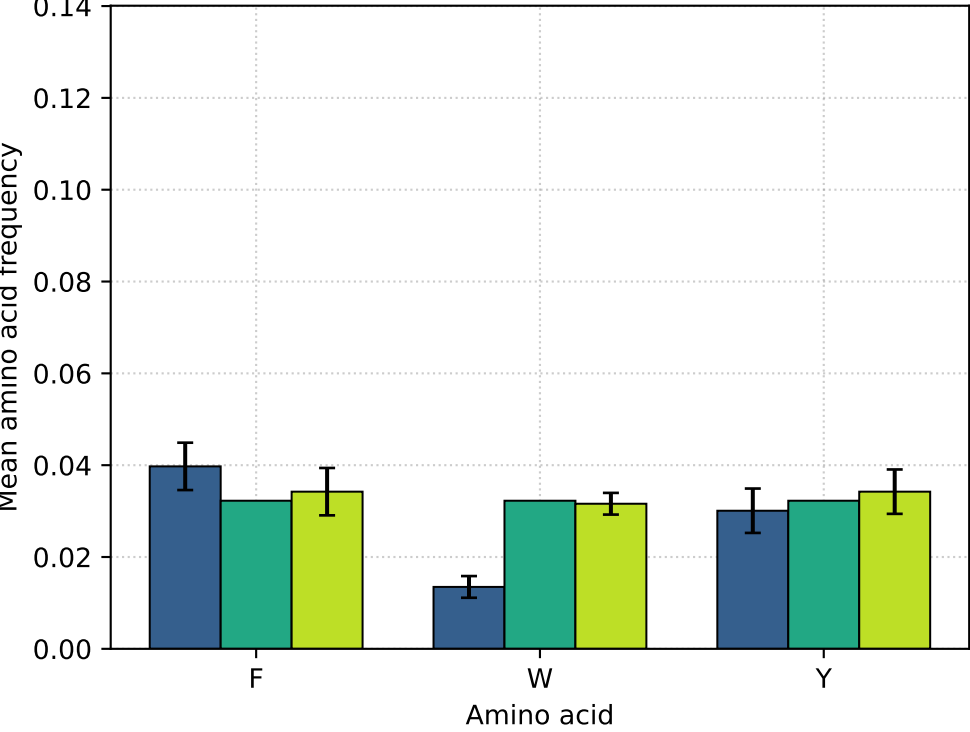
Aliphatic amino acids



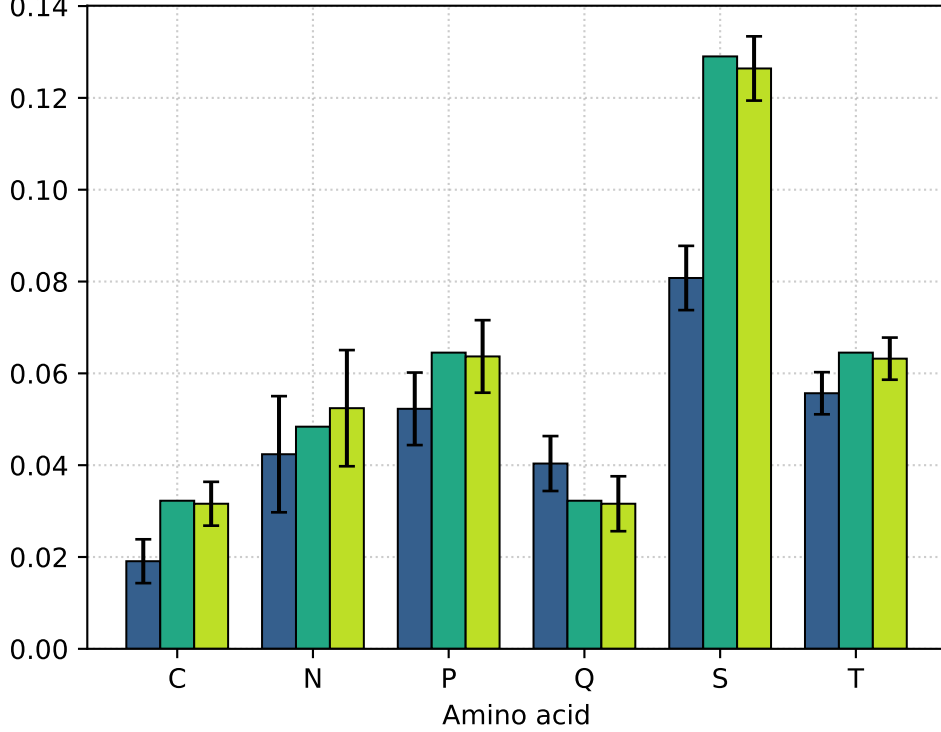
Charged amino acids



Aromatic amino acids



Uncharged amino acids



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