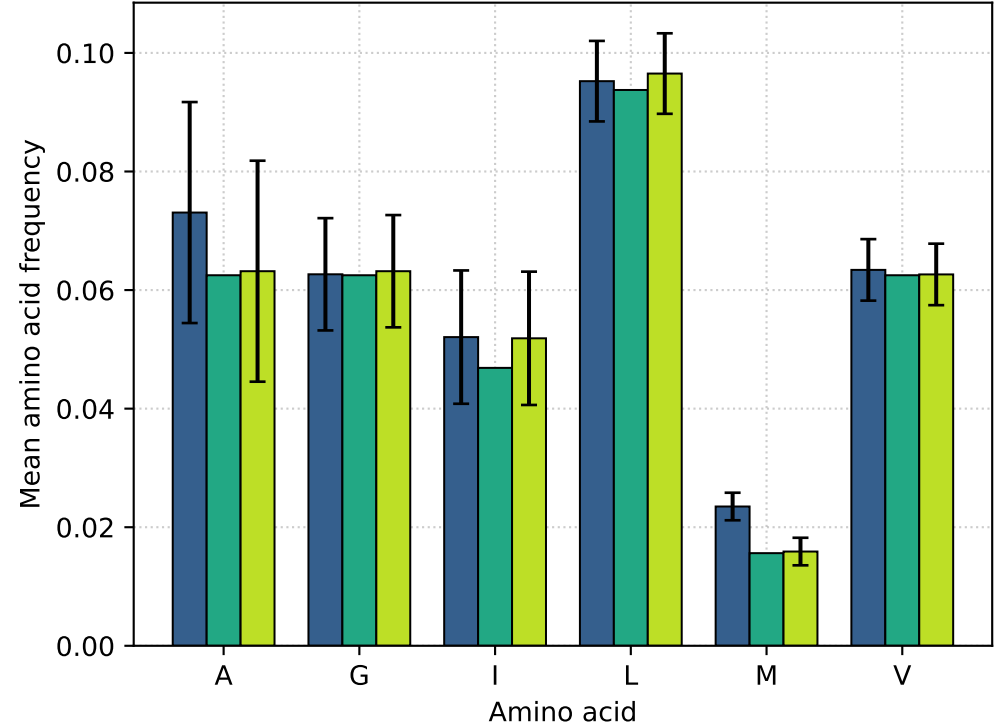
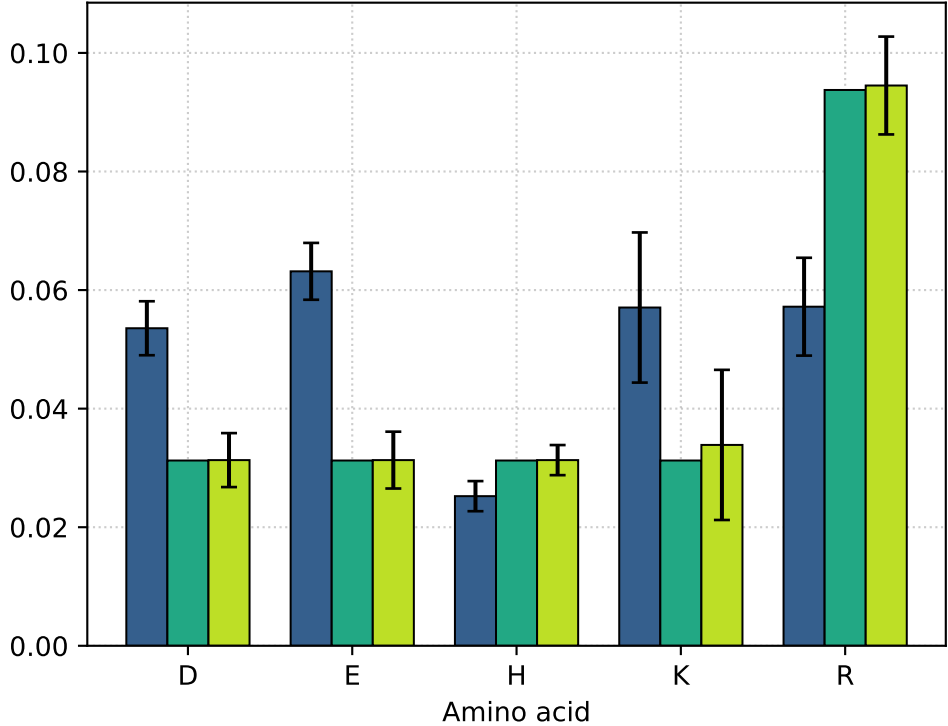


# Eukaryota - Mean proteome amino acid frequency for genetic code: Condyllostoma nuclear

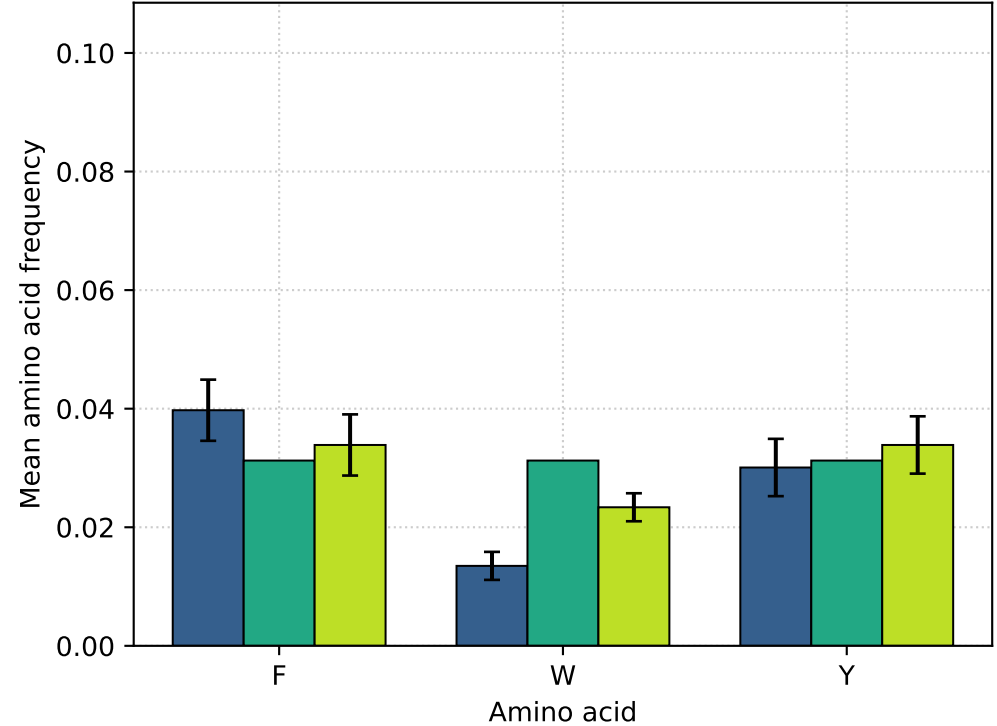
Aliphatic amino acids



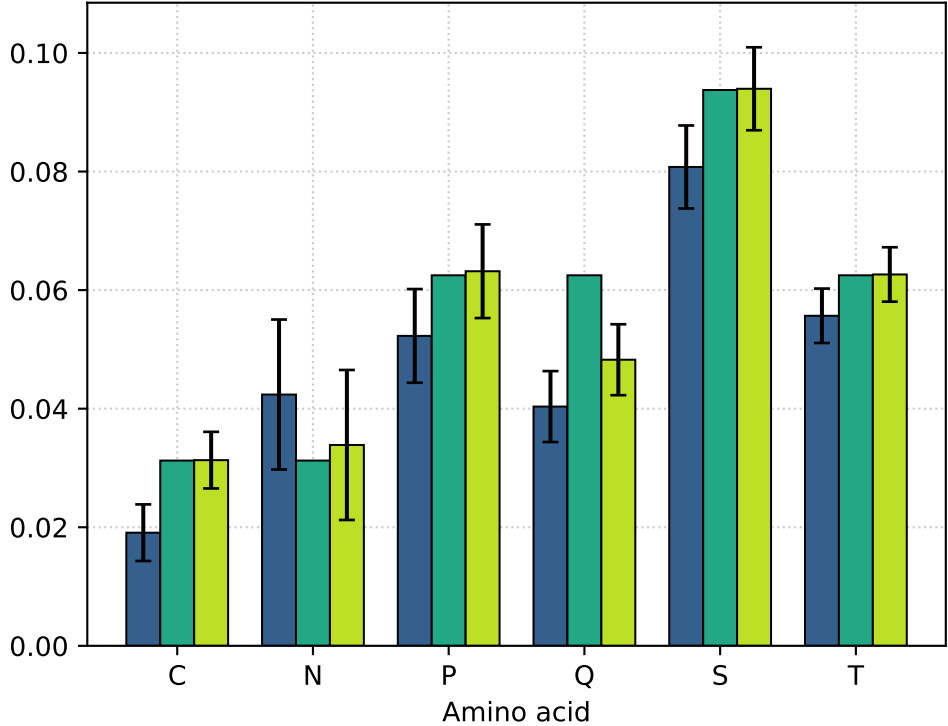
Charged amino acids



Aromatic amino acids



Uncharged amino acids



Observed  
Codon  
Codon+GC

Codon correlation:  
- Mean Pearson:  
- Coefficient: 0.69782  
- p-value: 1.68519e-02  
  
- Mean Spearman:  
- Coefficient: 0.67630  
- p-value: 2.28897e-02  
  
- Mean log-MSE: 0.36868

Codon + GC correlation:  
- Mean Pearson:  
- Coefficient: 0.77458  
- p-value: 2.71523e-04  
  
- Mean Spearman:  
- Coefficient: 0.74403  
- p-value: 8.84387e-04  
  
- Mean log-MSE: 0.29236