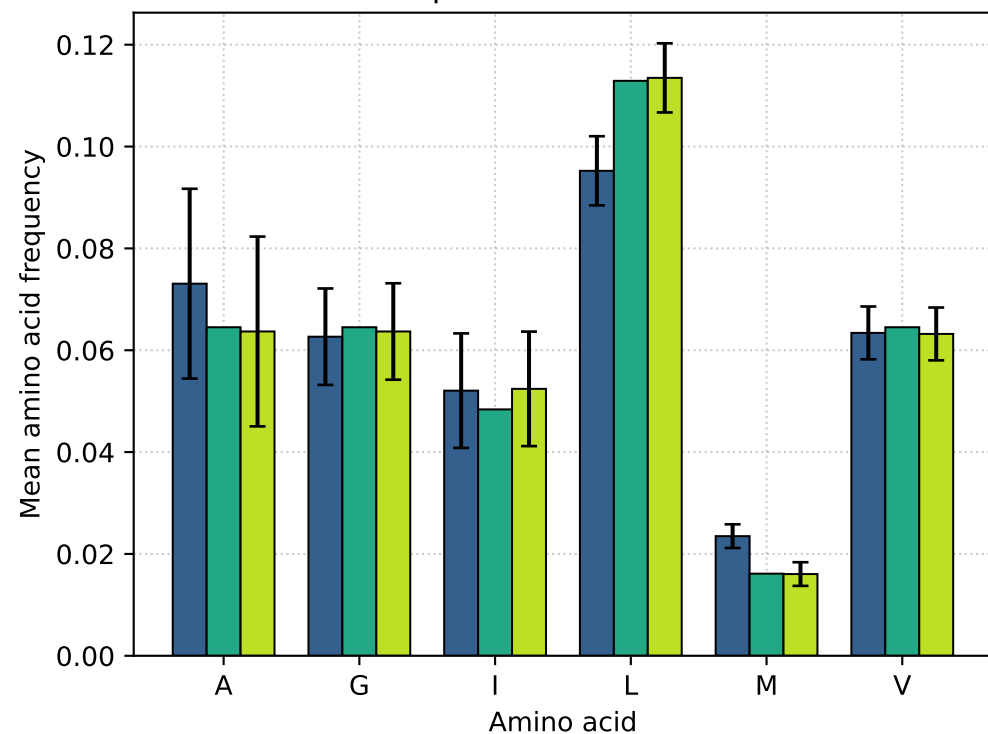
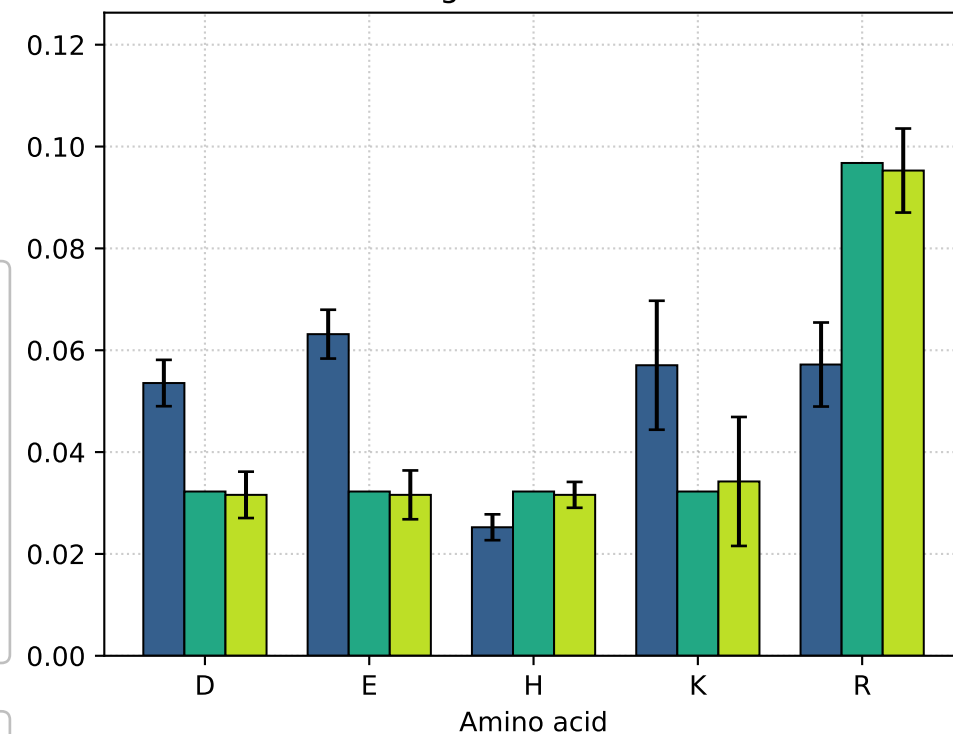


# Eukaryota - Mean proteome amino acid frequency for genetic code: Chlorophycean mitochondrial

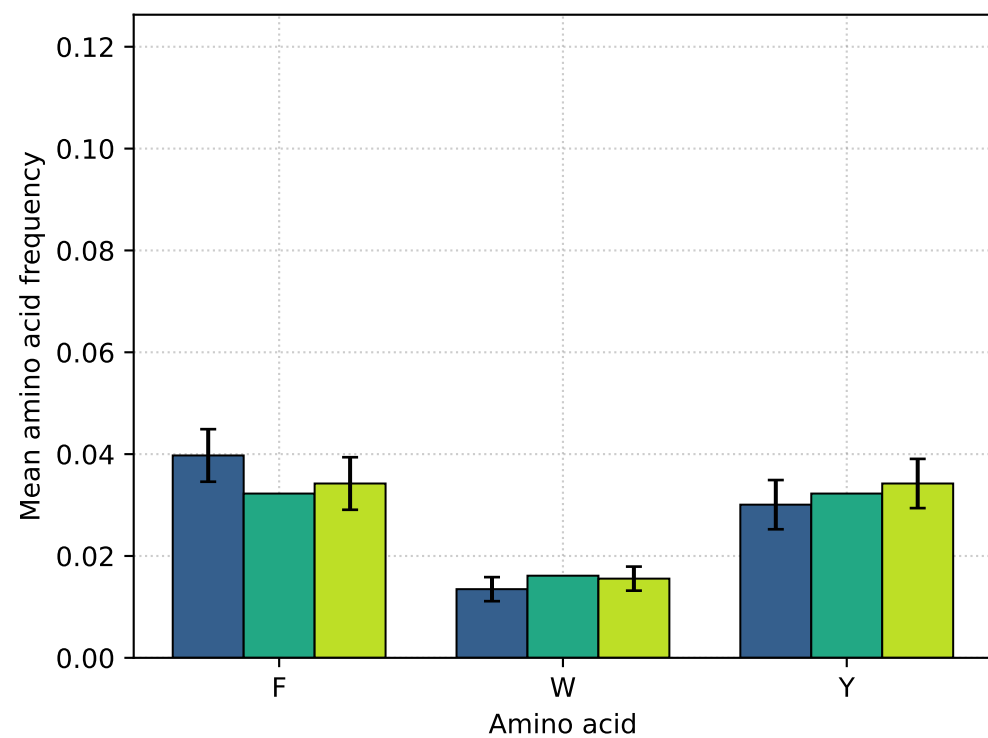
## Aliphatic amino acids



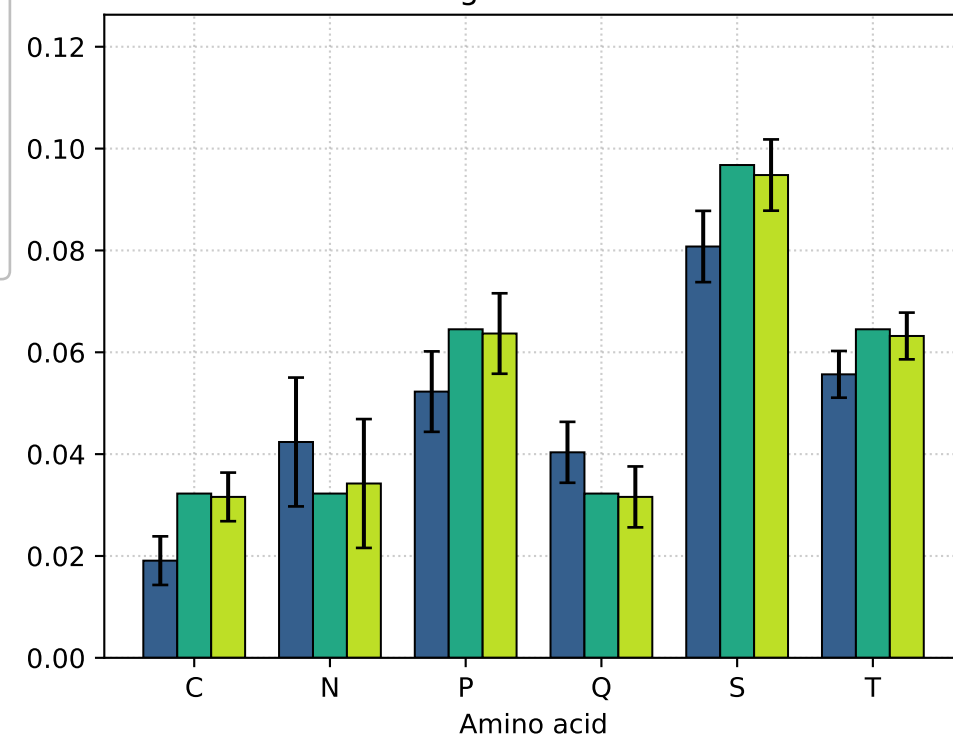
## Charged amino acids



## Aromatic amino acids



## Uncharged amino acids



Observed  
Codon  
Codon+GC

Codon correlation:  
- Mean Pearson:  
- Coefficient: 0.76725  
- p-value: 1.16905e-02  
  
- Mean Spearman:  
- Coefficient: 0.75916  
- p-value: 9.79261e-03  
  
- Mean log-MSE: 0.27737

Codon + GC correlation:  
- Mean Pearson:  
- Coefficient: 0.80385  
- p-value: 1.49875e-04  
  
- Mean Spearman:  
- Coefficient: 0.76920  
- p-value: 3.21168e-04  
  
- Mean log-MSE: 0.26460