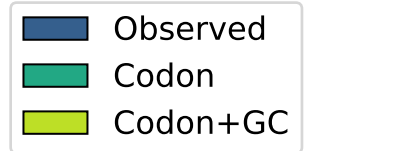
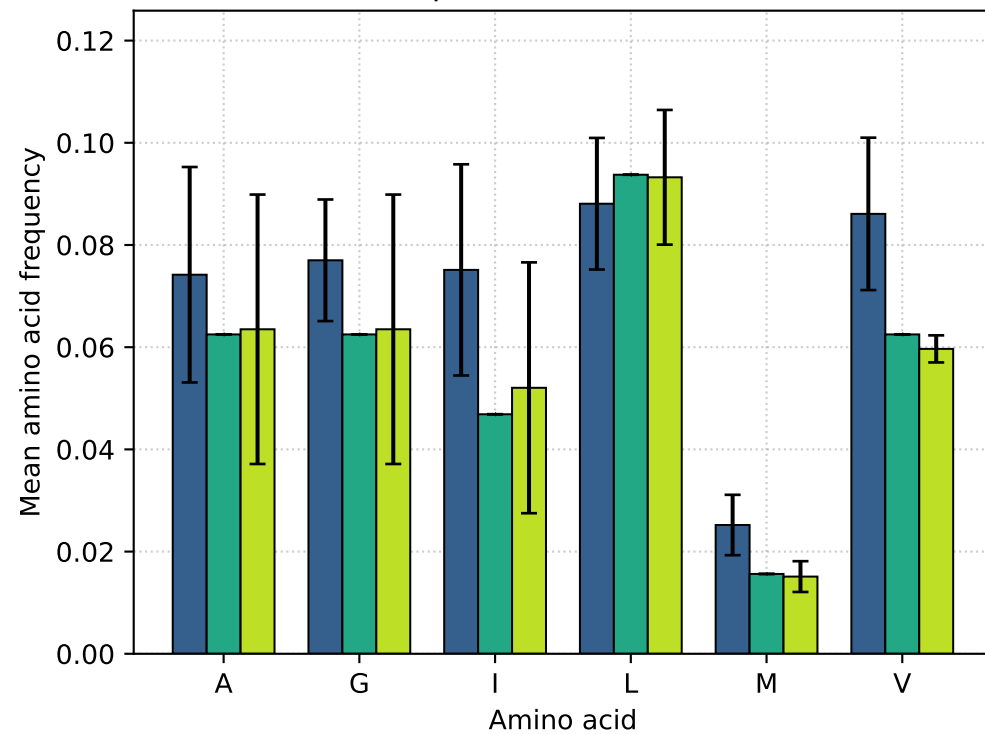


# Archaea - Mean proteome amino acid frequency for genetic code: Condyllostoma nuclear

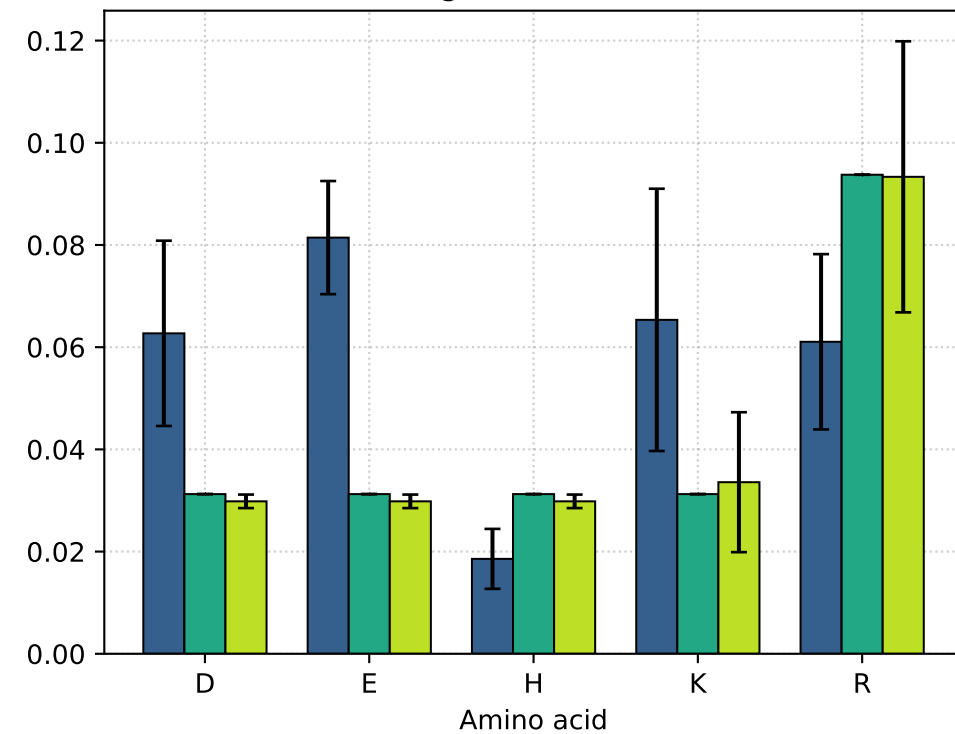
## Aliphatic amino acids



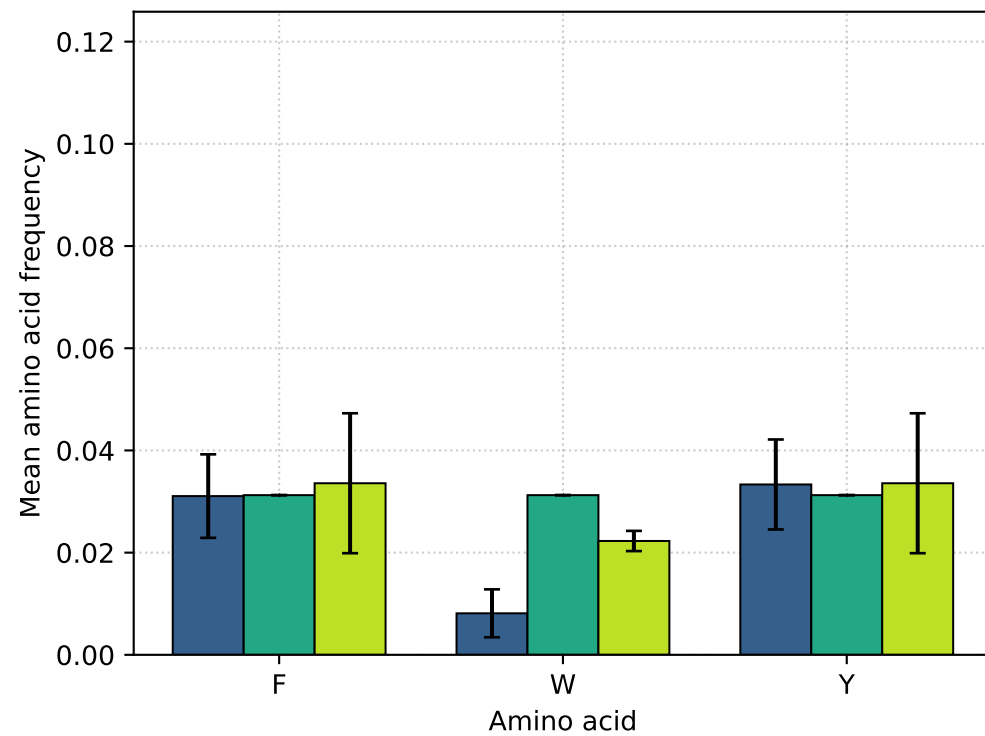
Codon correlation:  
- Mean Pearson:  
- Coefficient: 0.43134  
- p-value: 1.06759e-01  
  
- Mean Spearman:  
- Coefficient: 0.46342  
- p-value: 7.94714e-02

Codon + GC correlation:  
- Mean Pearson:  
- Coefficient: 0.52742  
- p-value: 4.29401e-02  
  
- Mean Spearman:  
- Coefficient: 0.53222  
- p-value: 3.63717e-02

## Charged amino acids



## Aromatic amino acids



## Uncharged amino acids

