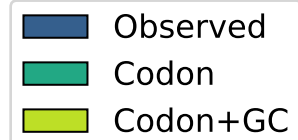
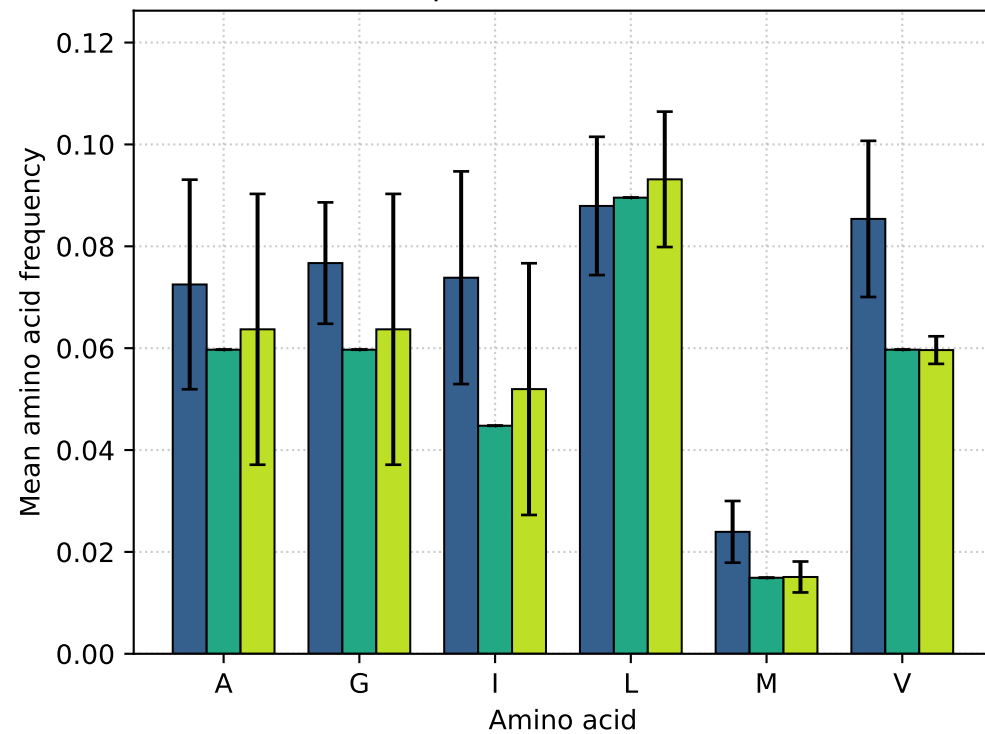


# Archaea - Mean proteomic amino acid frequency for genetic code: Condyllostoma nuclear

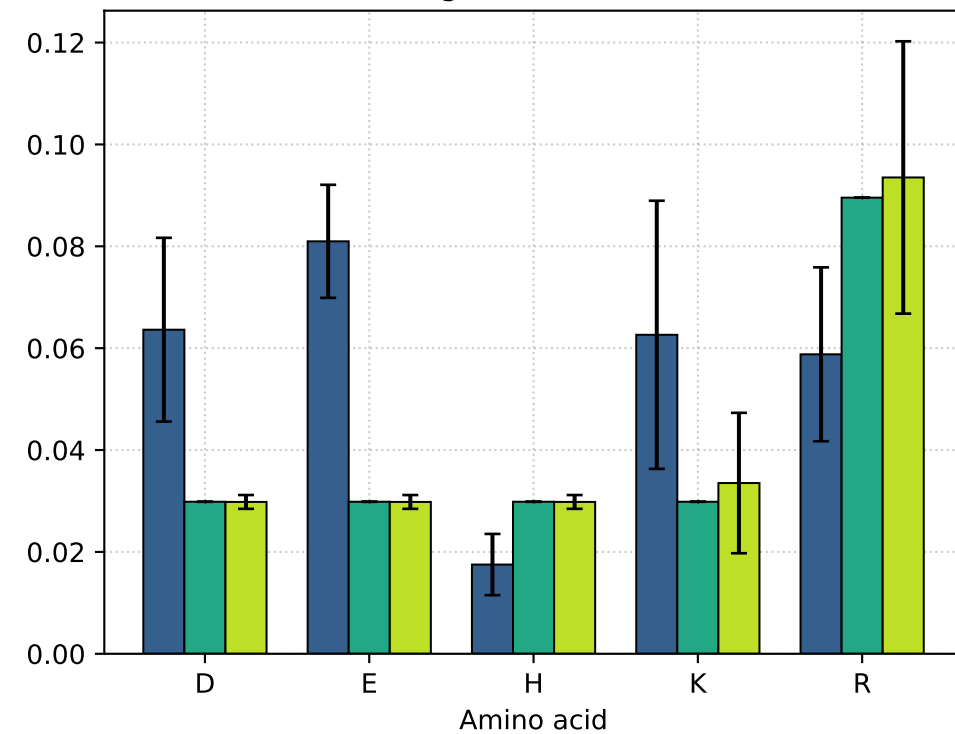
## Aliphatic amino acids



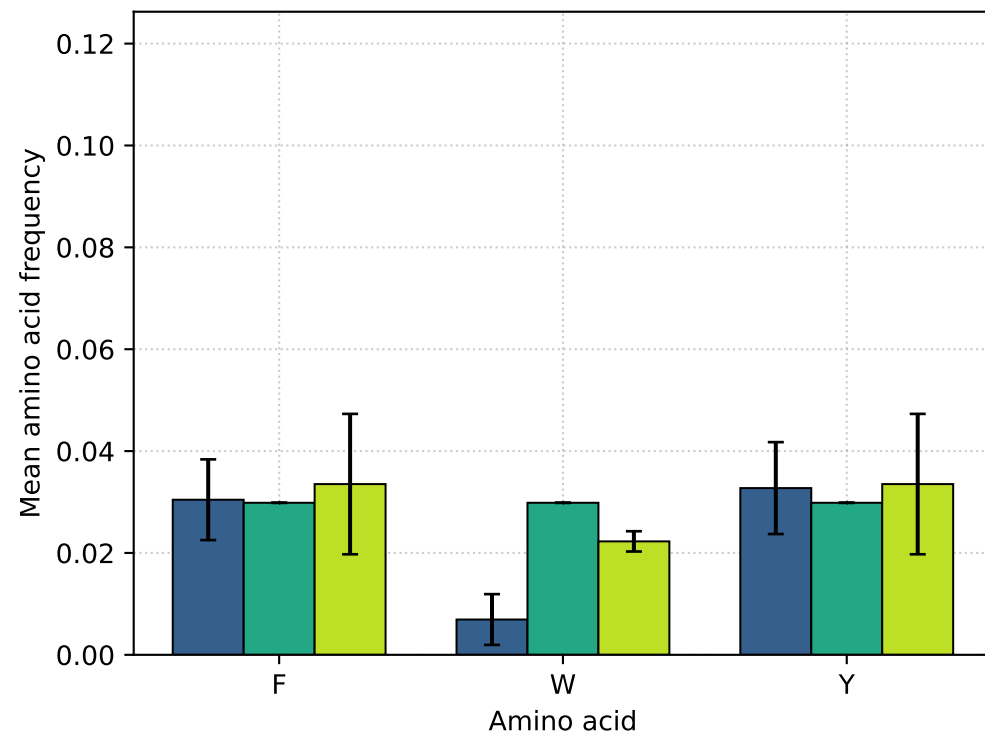
Codon correlation:  
- Pearson:  
- Coefficient: 0.428124  
- p-value: 0.107957  
  
- Spearman:  
- Coefficient: 0.460975  
- p-value: 0.077489

Codon + GC correlation:  
- Pearson:  
- Coefficient: 0.523146  
- p-value: 0.043585  
  
- Spearman:  
- Coefficient: 0.531630  
- p-value: 0.034926

## Charged amino acids



## Aromatic amino acids



## Uncharged amino acids

