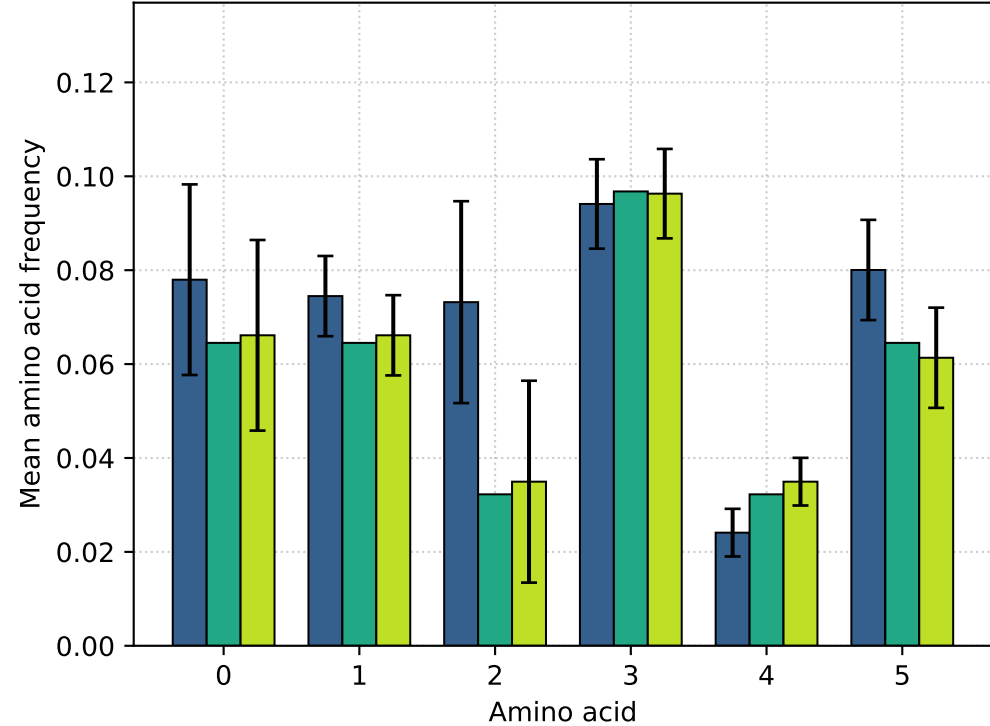
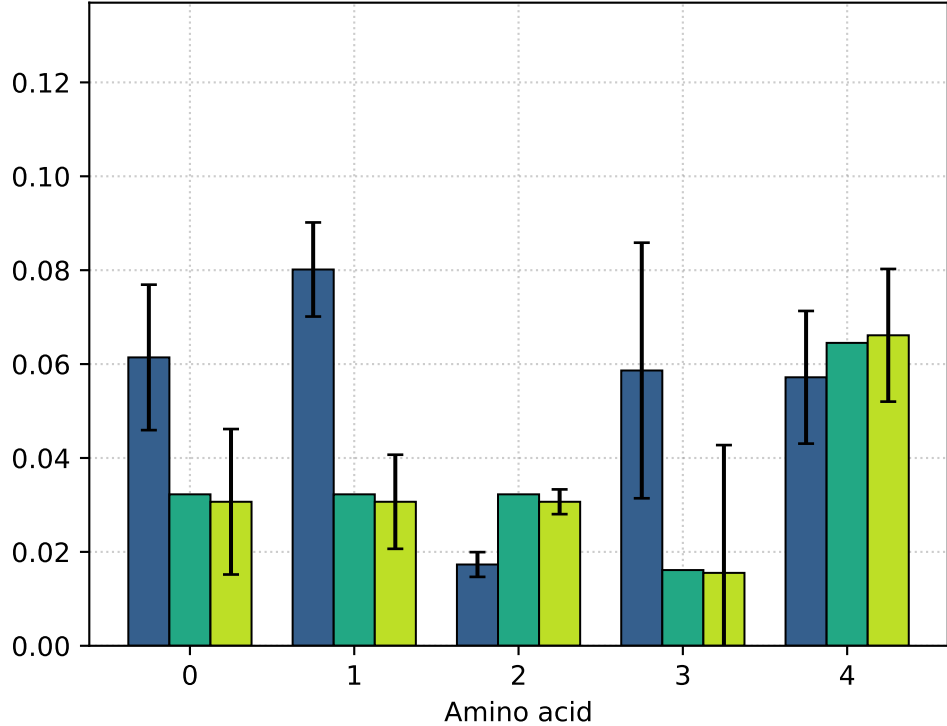


# Archaea - Mean proteome amino acid frequency for genetic code: Trematode mitochondrial

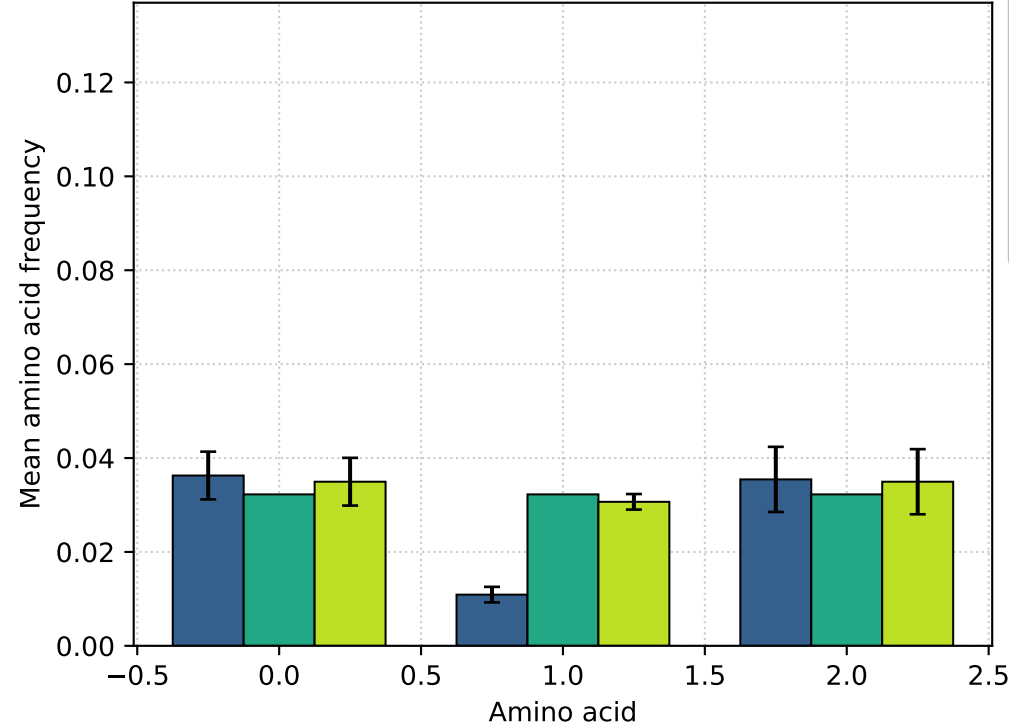
Aliphatic amino acids



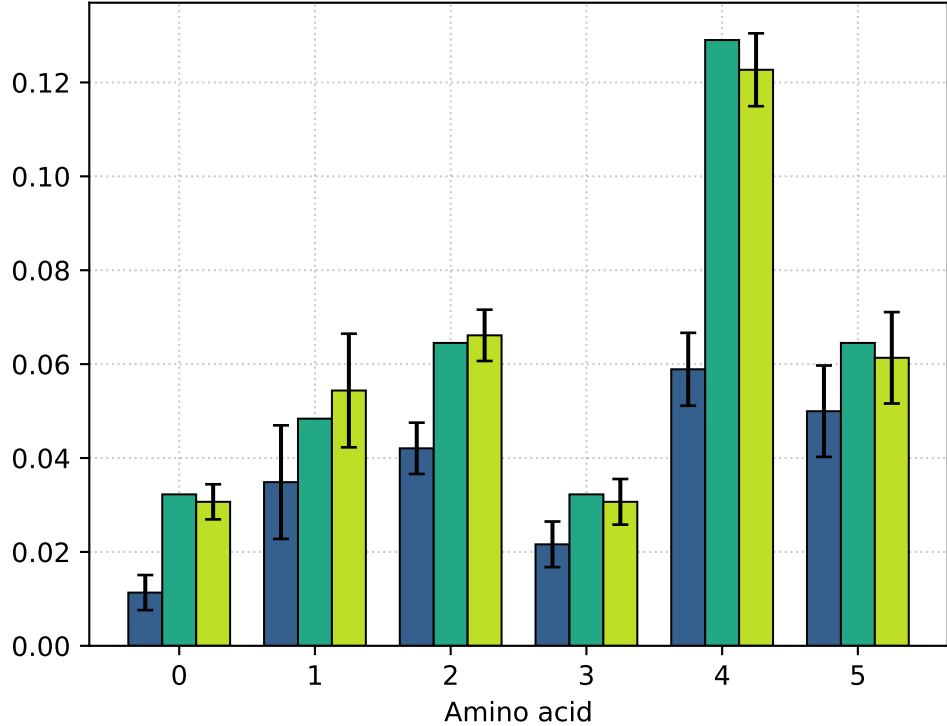
Charged amino acids



Aromatic amino acids



Uncharged amino acids



Codon correlation:  
- Mean Pearson:  
- Coefficient: 0.41092  
- p-value: 1.29383e-01  
  
- Mean Spearman:  
- Coefficient: 0.44434  
- p-value: 1.17167e-01  
  
- Mean log-MSE: 0.88487

Codon + GC correlation:  
- Mean Pearson:  
- Coefficient: 0.45986  
- p-value: 8.61822e-02  
  
- Mean Spearman:  
- Coefficient: 0.42281  
- p-value: 1.30417e-01  
  
- Mean log-MSE: 0.91511