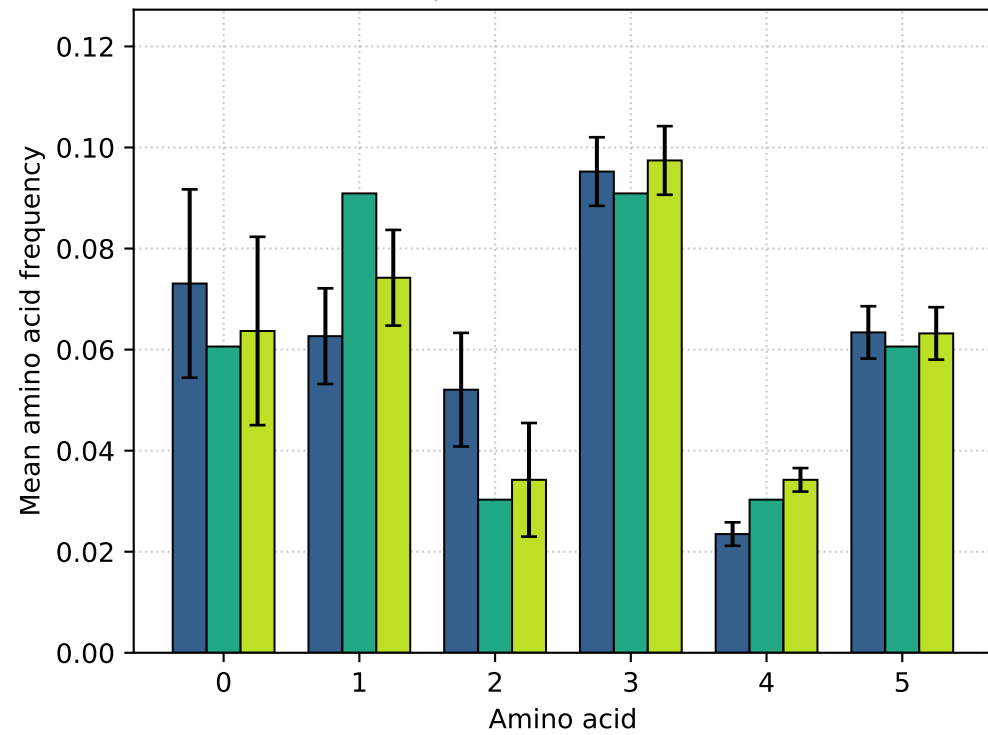
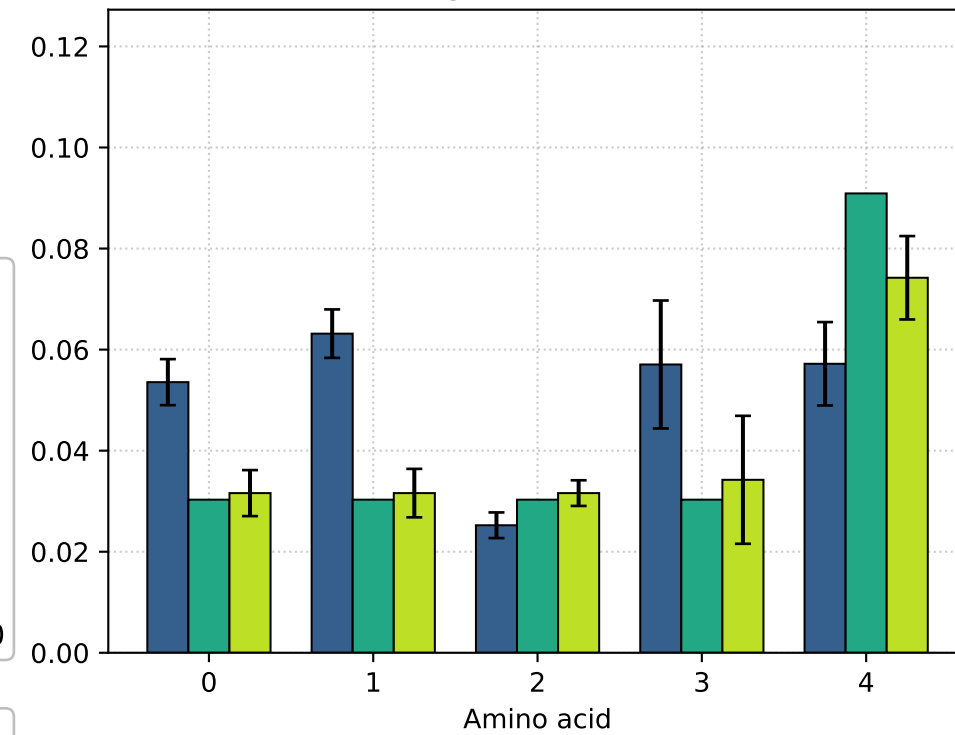


# Eukaryota - Mean proteome amino acid frequency for genetic code: Ascidian mitochondrial

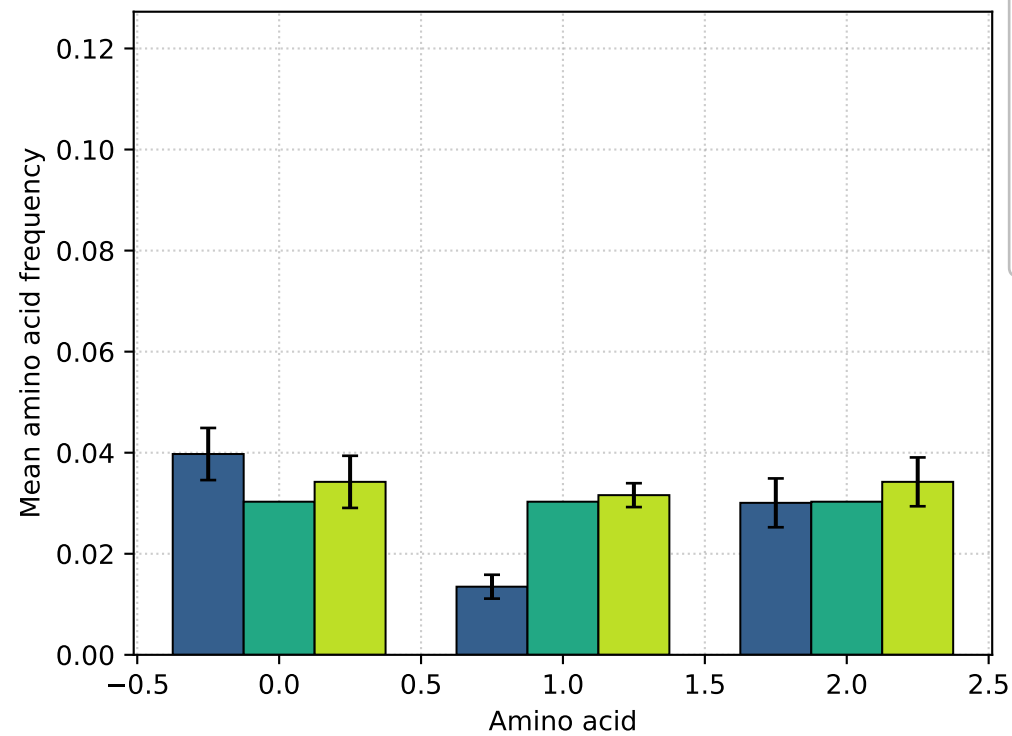
## Aliphatic amino acids



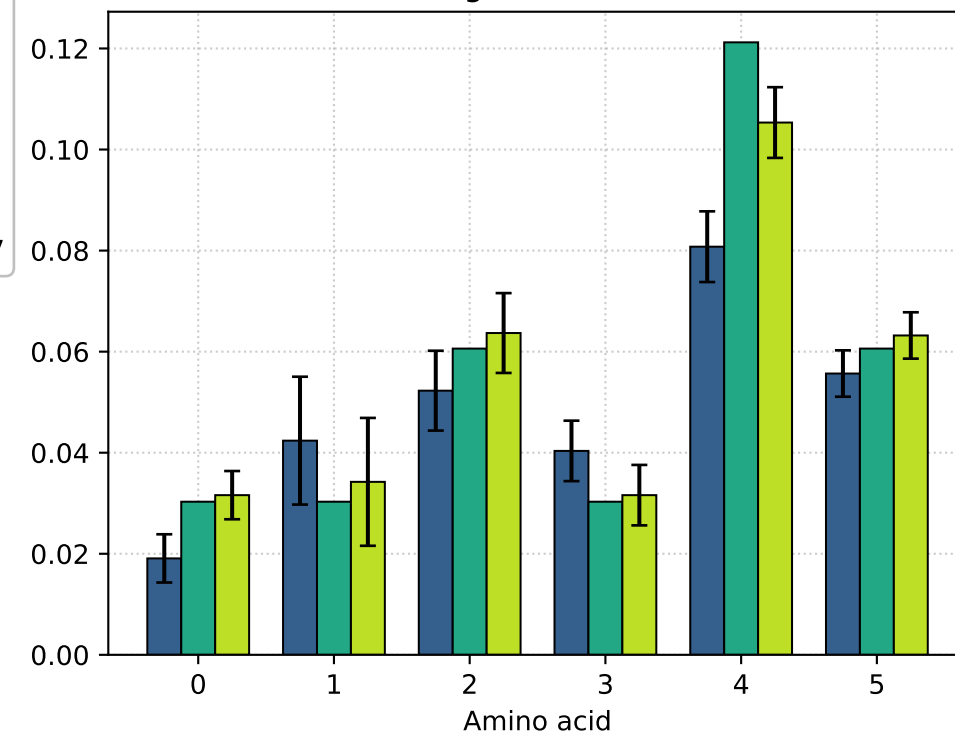
## Charged amino acids



## Aromatic amino acids



## Uncharged amino acids



Codon correlation:  
- Mean Pearson:  
- Coefficient: 0.67423  
- p-value: 2.13380e-02

- Mean Spearman:  
- Coefficient: 0.68385  
- p-value: 3.23269e-02  
- Mean log-MSE: 0.40520

Codon + GC correlation:  
- Mean Pearson:  
- Coefficient: 0.76056  
- p-value: 4.93064e-04

- Mean Spearman:  
- Coefficient: 0.68088  
- p-value: 2.82594e-03  
- Mean log-MSE: 0.34617