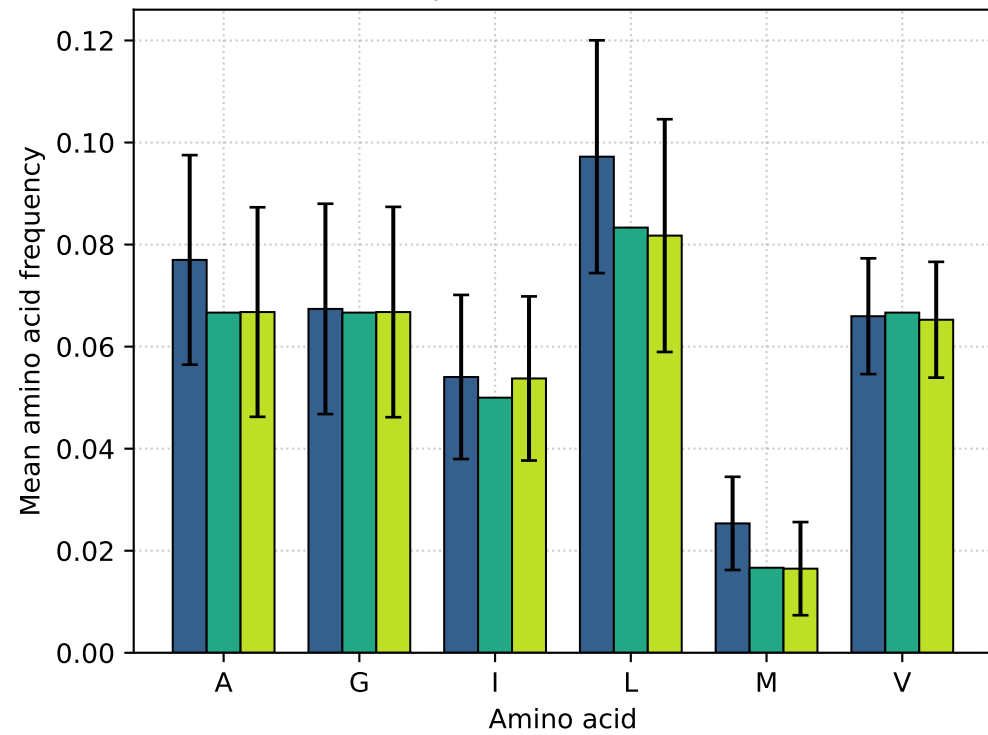


# Eukaryota - Mean proteome amino acid frequency for genetic code: Thraustochytrium mitochondrial

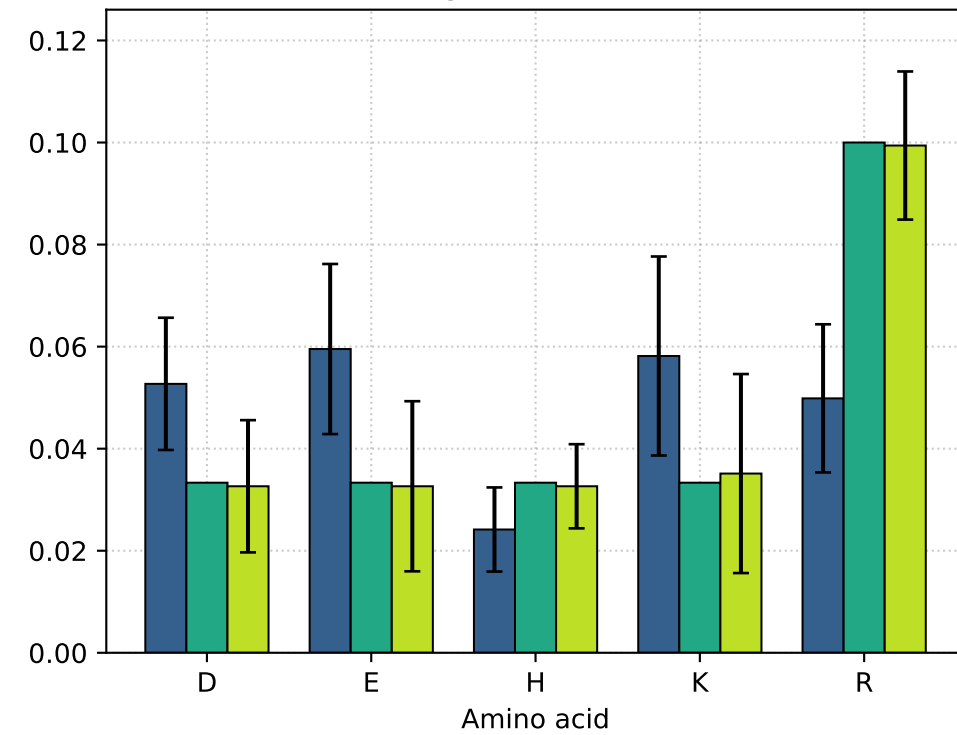
## Aliphatic amino acids



Observed  
Codon  
Codon+GC

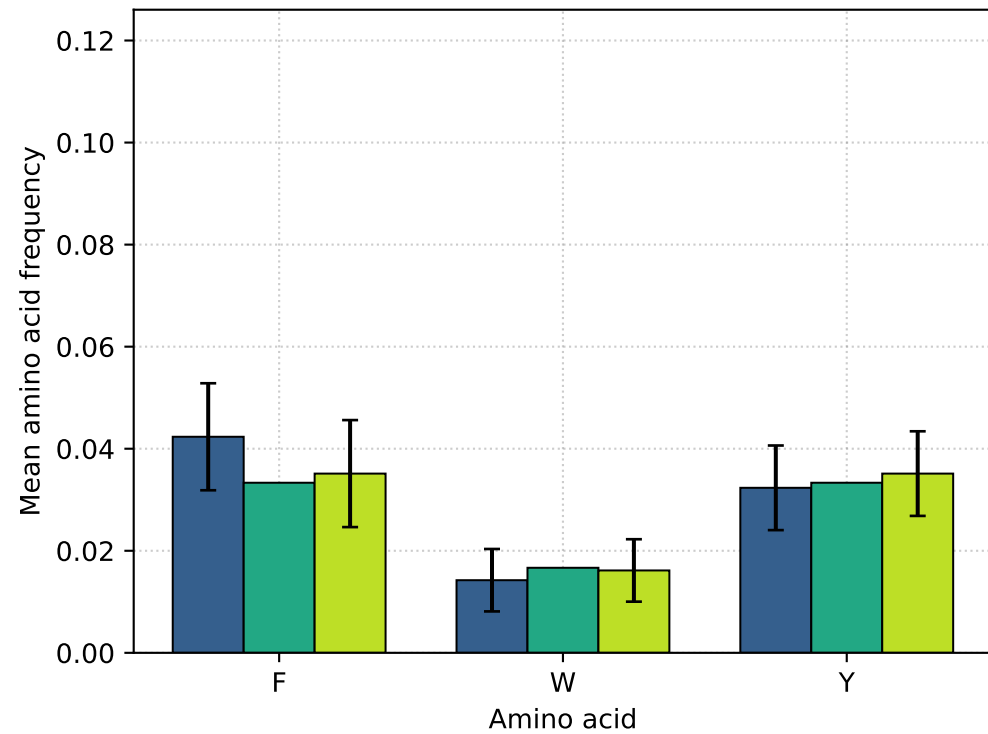
Codon correlation:  
- Mean Pearson:  
- Coefficient: 0.60785  
- p-value: 3.15904e-02  
  
- Mean Spearman:  
- Coefficient: 0.65222  
- p-value: 1.92415e-02

## Charged amino acids



Codon + GC correlation:  
- Mean Pearson:  
- Coefficient: 0.63546  
- p-value: 1.76158e-02  
  
- Mean Spearman:  
- Coefficient: 0.65361  
- p-value: 1.27584e-02

## Aromatic amino acids



## Uncharged amino acids

