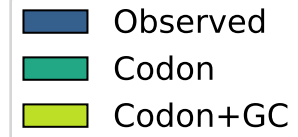
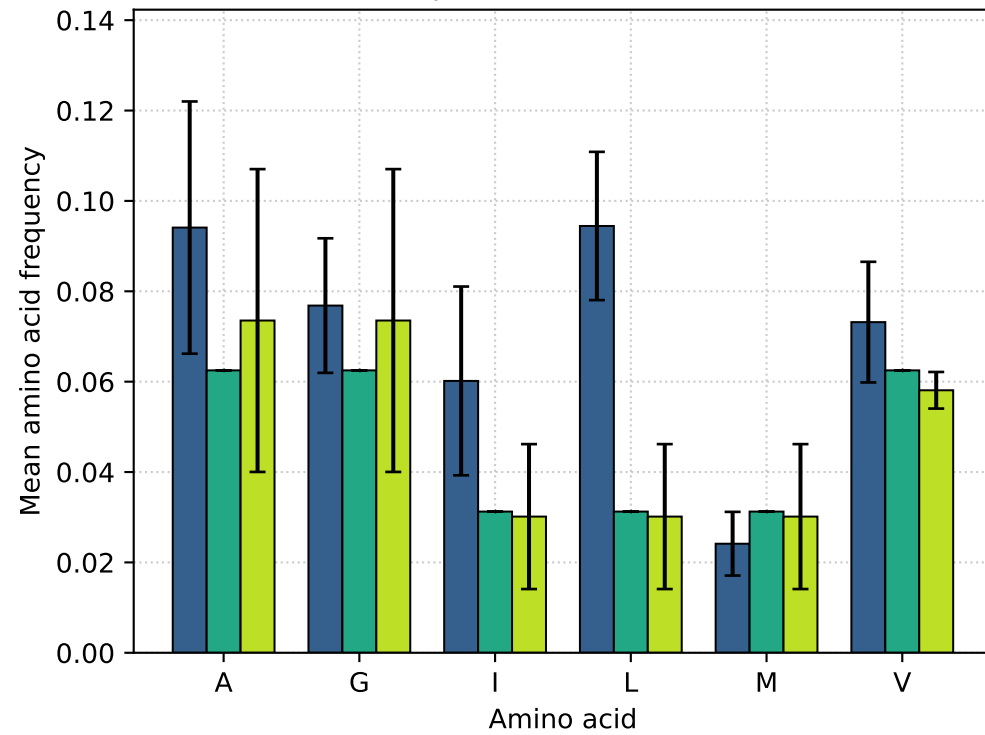


# Bacteria - Mean proteomic amino acid frequency for genetic code: Yeast mitochondrial

## Aliphatic amino acids



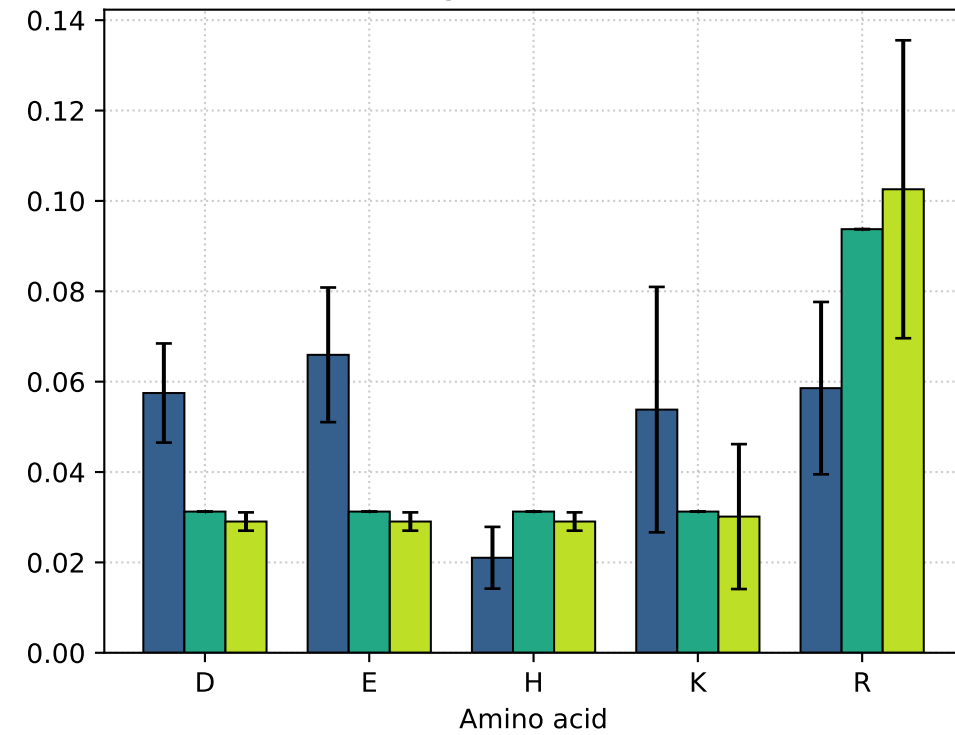
**Codon correlation:**

- Pearson:  
- Coefficient: 0.256016  
- p-value: 0.320838
- Spearman:  
- Coefficient: 0.377988  
- p-value: 0.188694

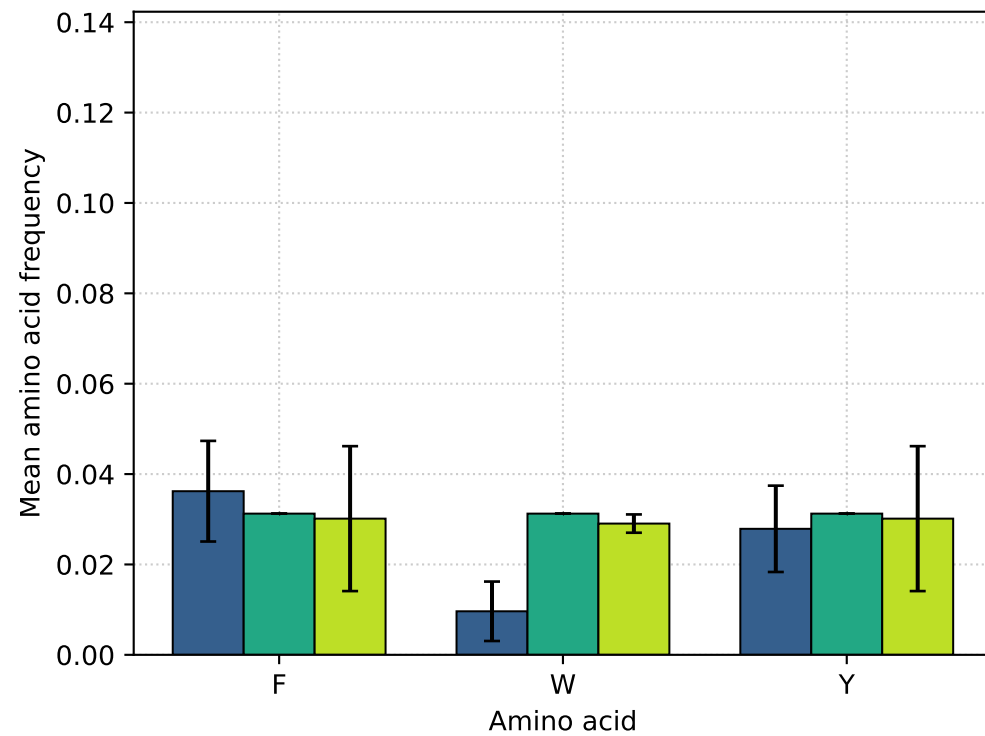
**Codon + GC correlation:**

- Pearson:  
- Coefficient: 0.363083  
- p-value: 0.204107
- Spearman:  
- Coefficient: 0.389527  
- p-value: 0.133509

## Charged amino acids



## Aromatic amino acids



## Uncharged amino acids

