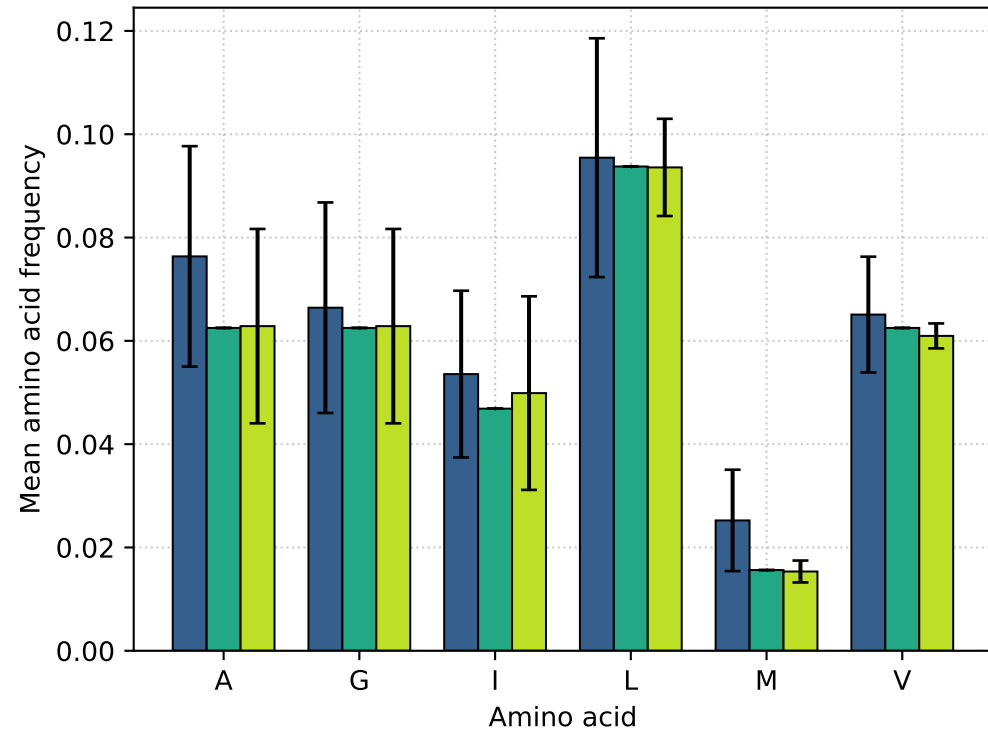


# Eukaryota - Mean proteomic amino acid frequency for genetic code: Mold-protzoan-coelenterate mitochondrial and mycoplasma spiroplasma

## Aliphatic amino acids



Observed  
Codon  
Codon+GC

### Codon correlation:

- Pearson:  
- Coefficient: 0.610781  
- p-value: 0.033931

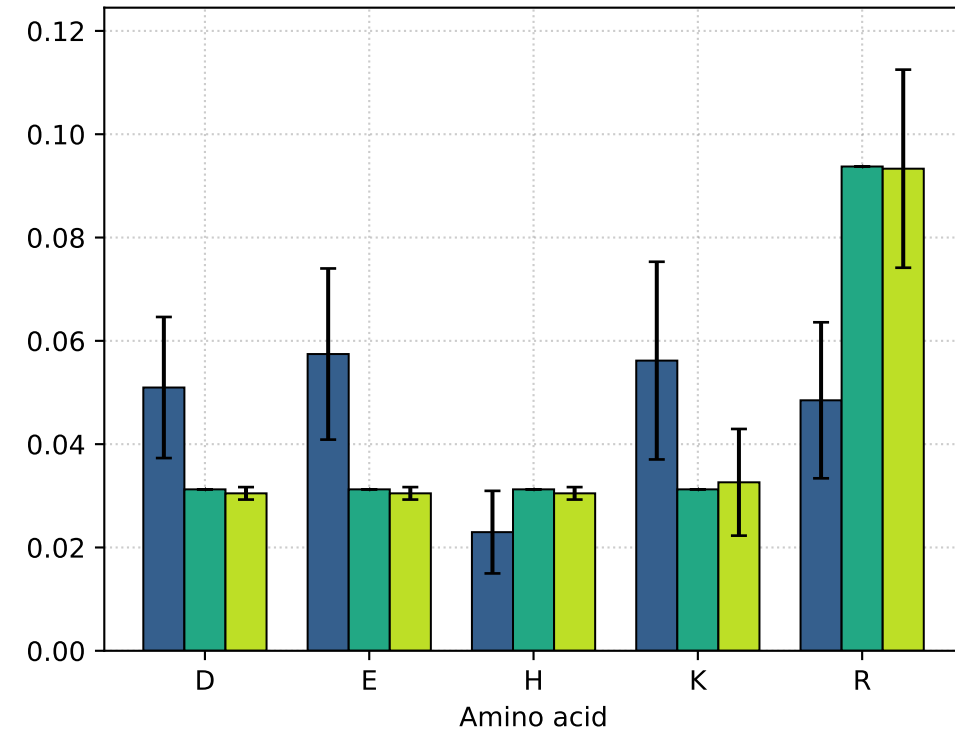
- Spearman:  
- Coefficient: 0.627409  
- p-value: 0.028461

### Codon + GC correlation:

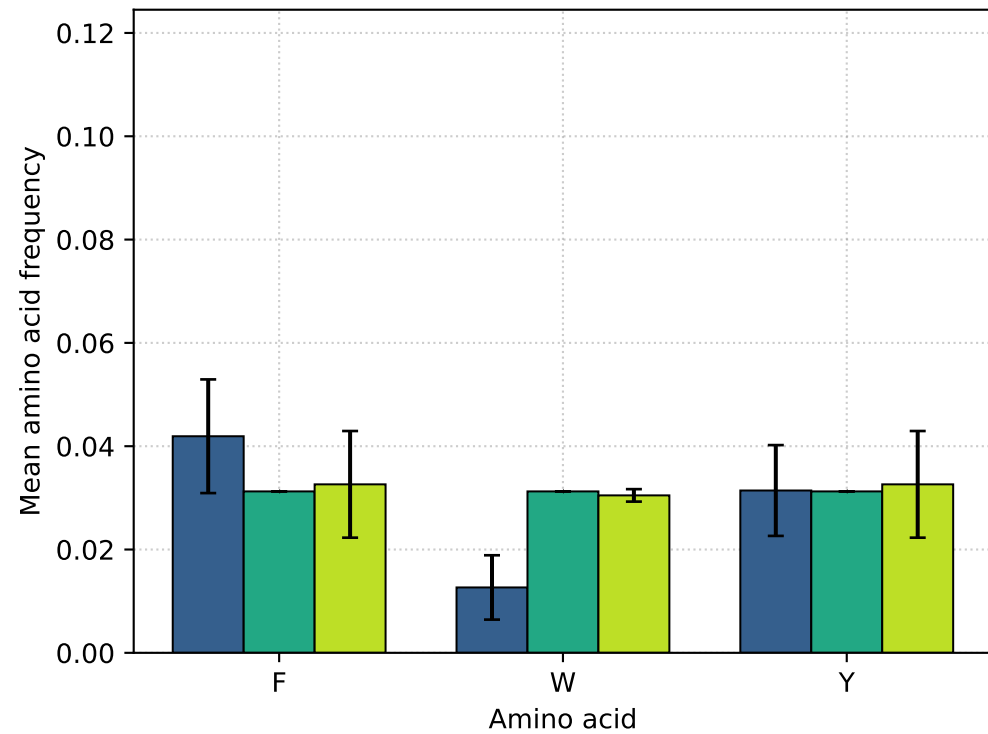
- Pearson:  
- Coefficient: 0.640256  
- p-value: 0.018263

- Spearman:  
- Coefficient: 0.617819  
- p-value: 0.020389

## Charged amino acids



## Aromatic amino acids



## Uncharged amino acids

