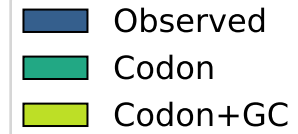
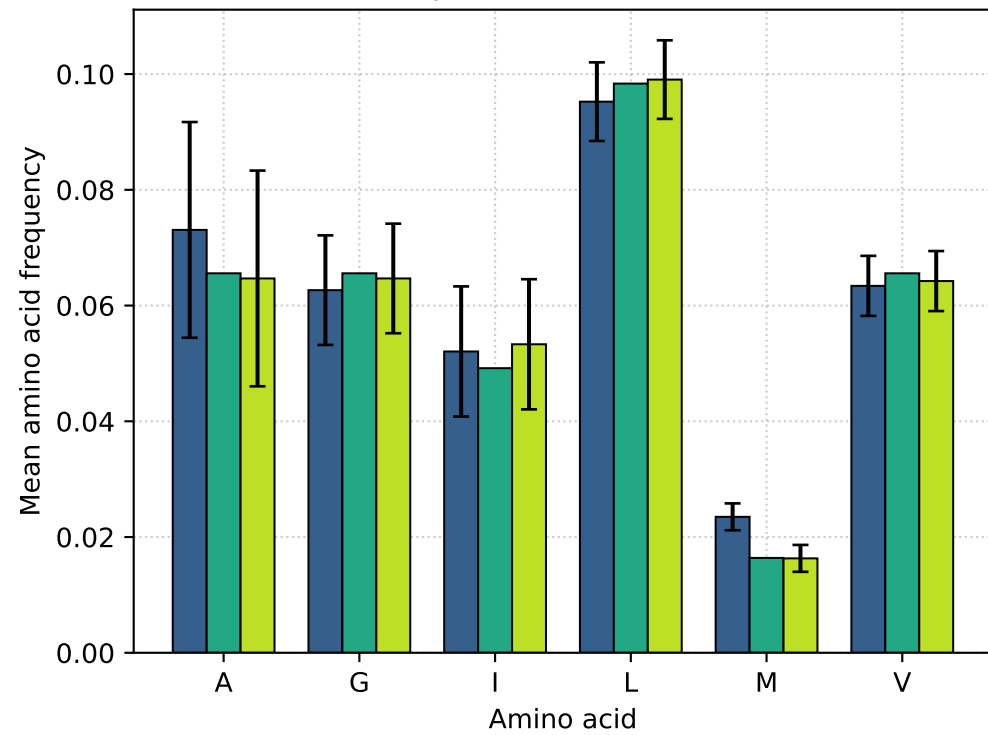


# Eukaryota - Mean proteome amino acid frequency for genetic code: Bacterial-archaeal-plant plastid

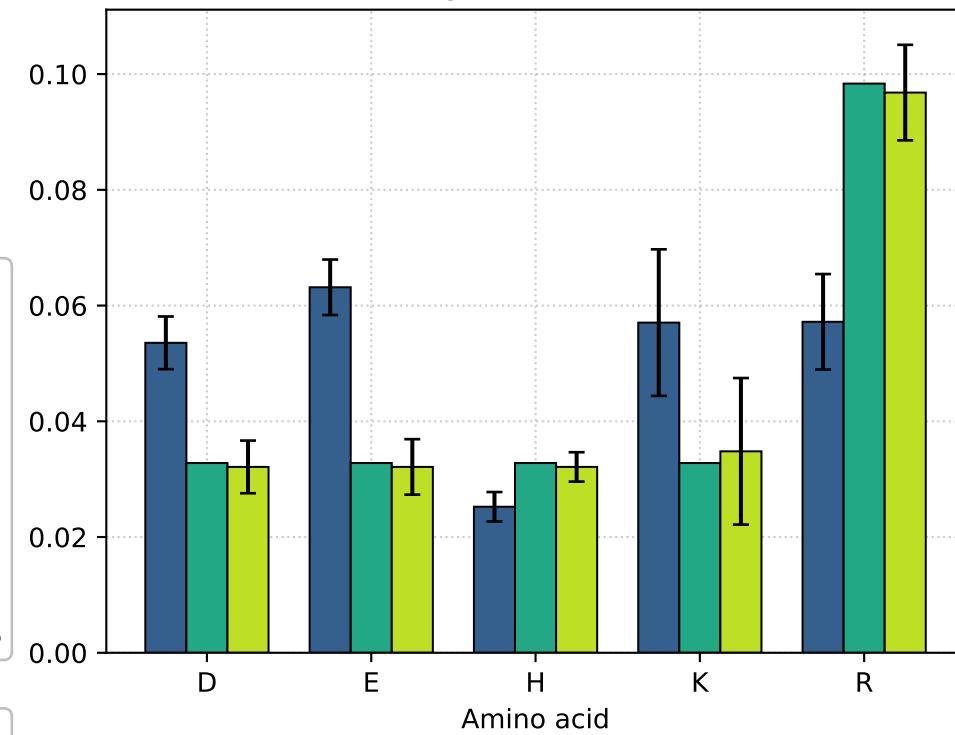
## Aliphatic amino acids



*Codon correlation:*

- Mean Pearson:  
- Coefficient: 0.75245  
- p-value: 1.38887e-02
- Mean Spearman:  
- Coefficient: 0.75366  
- p-value: 1.05887e-02
- Mean log-MSE: 0.27275

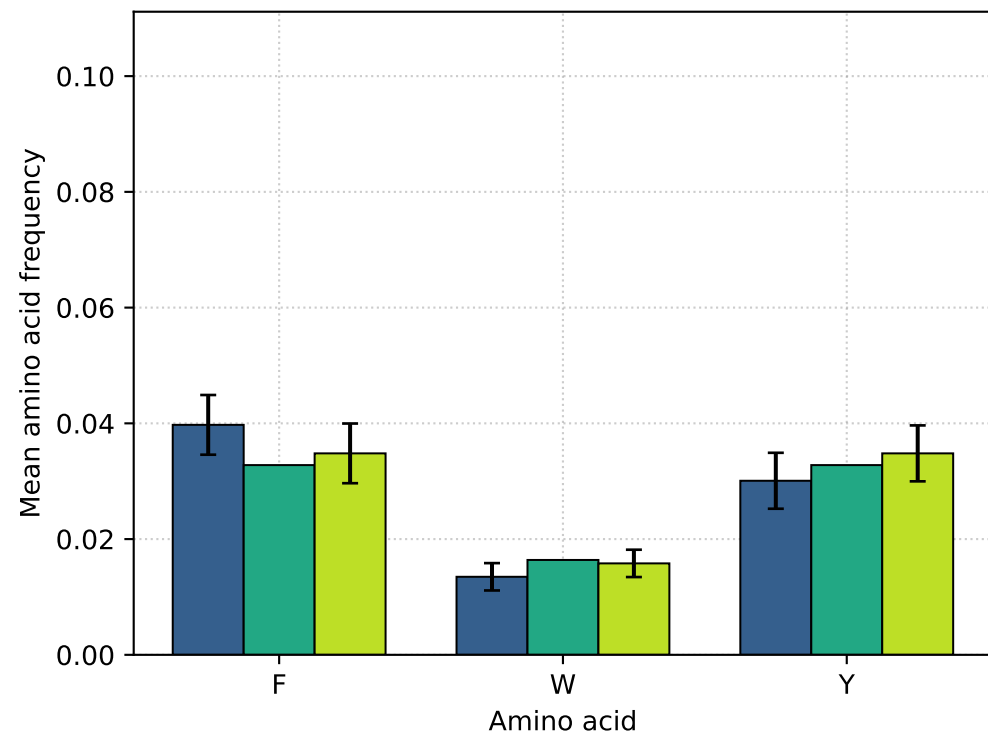
## Charged amino acids



*Codon + GC correlation:*

- Mean Pearson:  
- Coefficient: 0.79252  
- p-value: 1.62267e-04
- Mean Spearman:  
- Coefficient: 0.76515  
- p-value: 3.36228e-04
- Mean log-MSE: 0.25888

## Aromatic amino acids



## Uncharged amino acids

