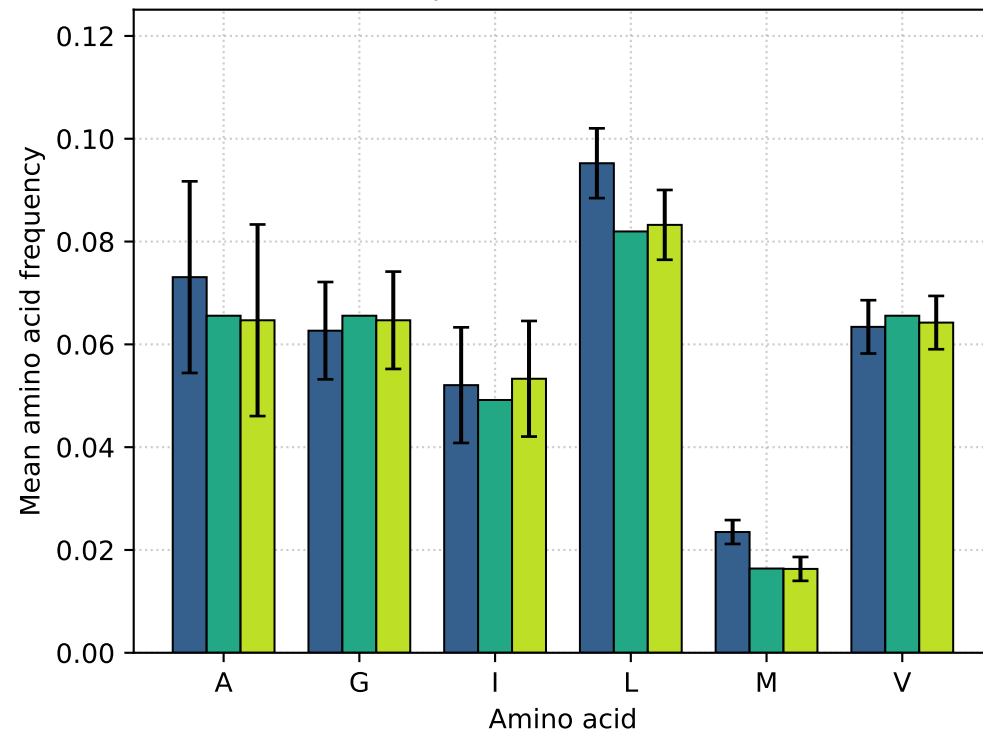
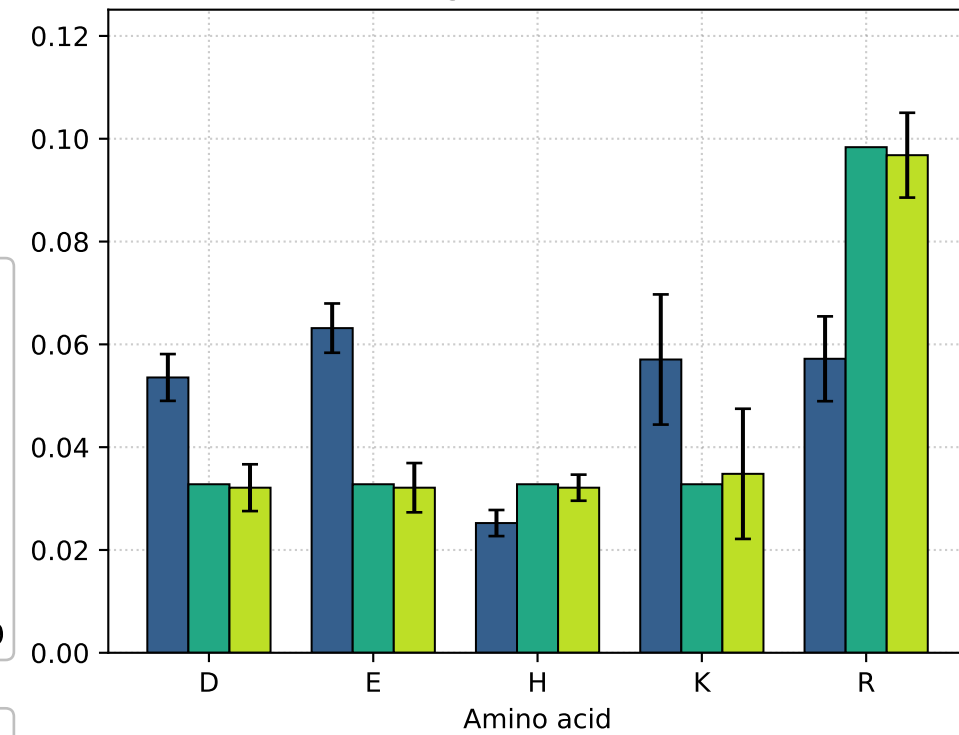


# Eukaryota - Mean proteome amino acid frequency for genetic code: Alternative yeast

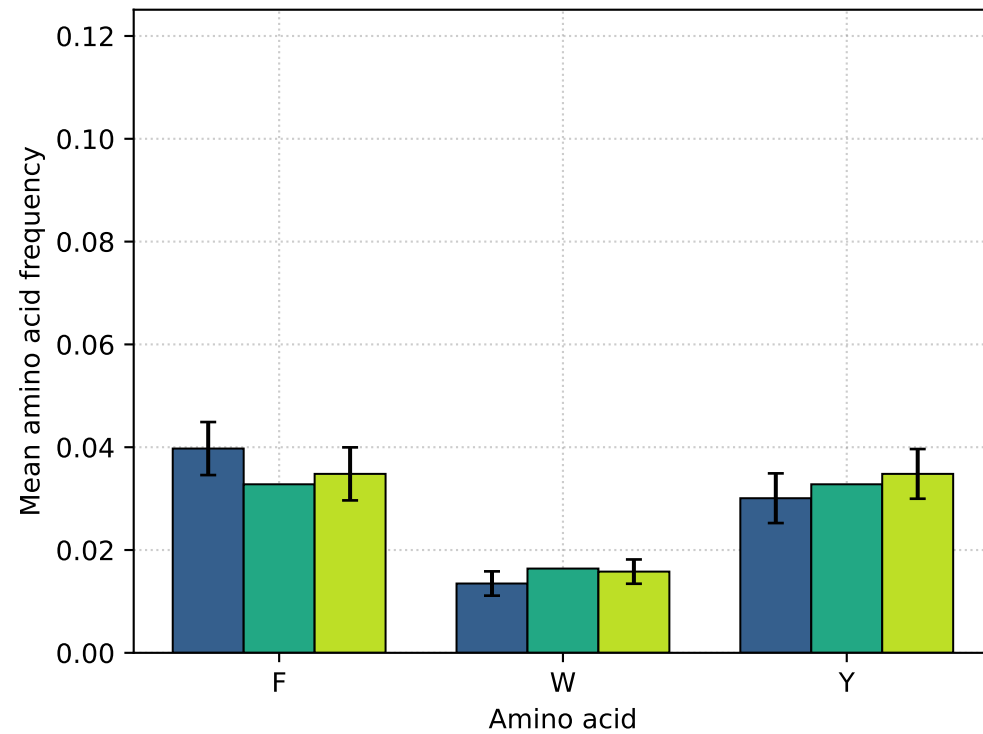
## Aliphatic amino acids



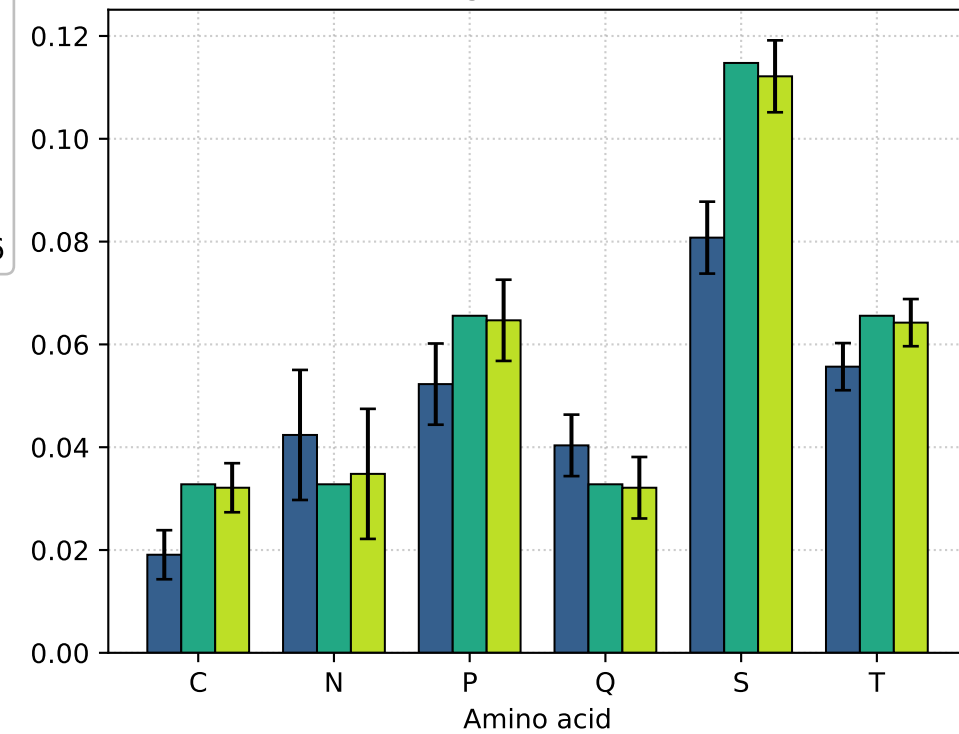
## Charged amino acids



## Aromatic amino acids



## Uncharged amino acids



Observed  
Codon  
Codon+GC

### Codon correlation:

- Mean Pearson:
- Coefficient: 0.71663
- p-value: 1.60912e-02

- Mean Spearman:
- Coefficient: 0.75044
- p-value: 1.09036e-02

- Mean log-MSE: 0.28379

### Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.76120
- p-value: 3.22818e-04

- Mean Spearman:
- Coefficient: 0.76227
- p-value: 3.62235e-04

- Mean log-MSE: 0.27036