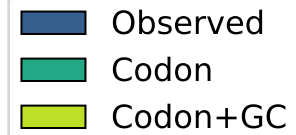
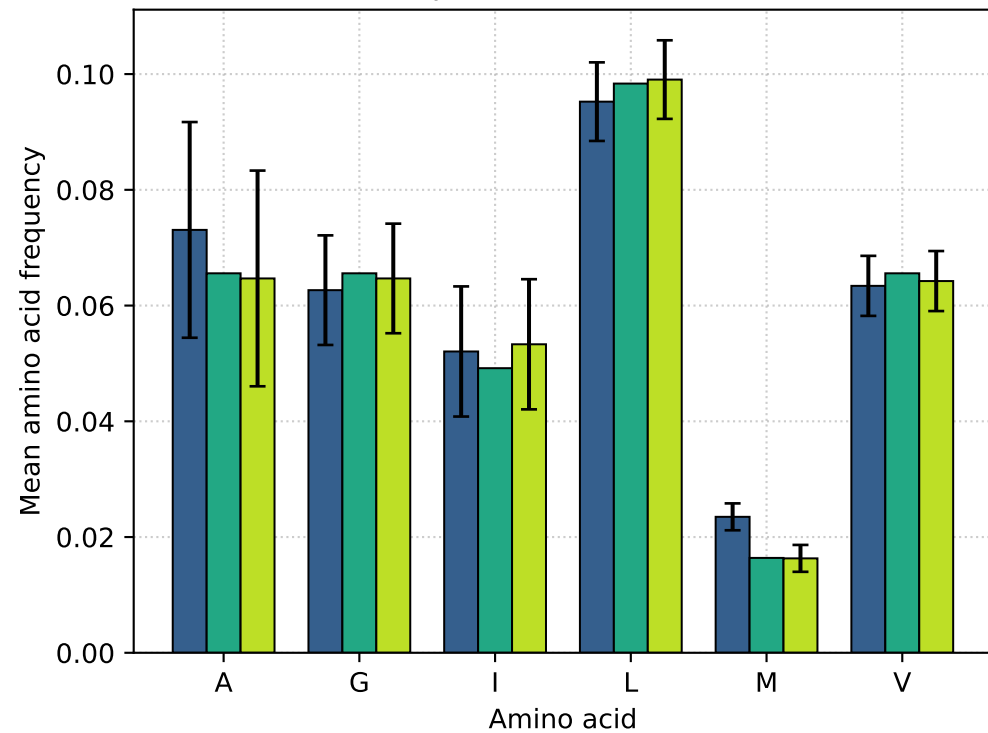


Eukaryota - Mean proteome amino acid frequency for genetic code: Standard

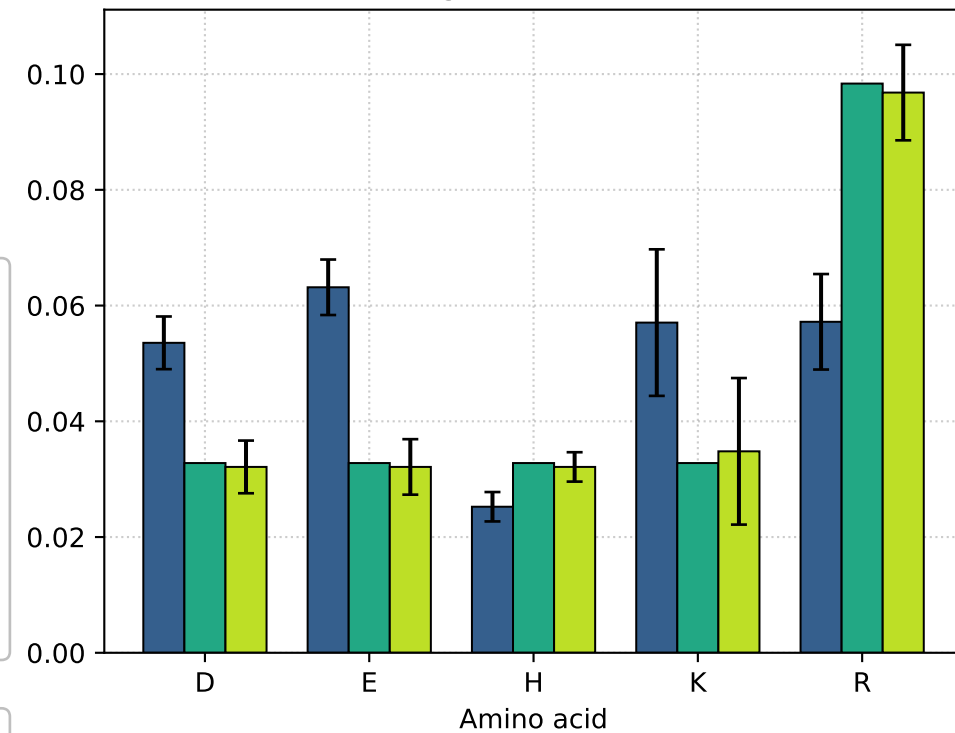
Aliphatic amino acids



Codon correlation:

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

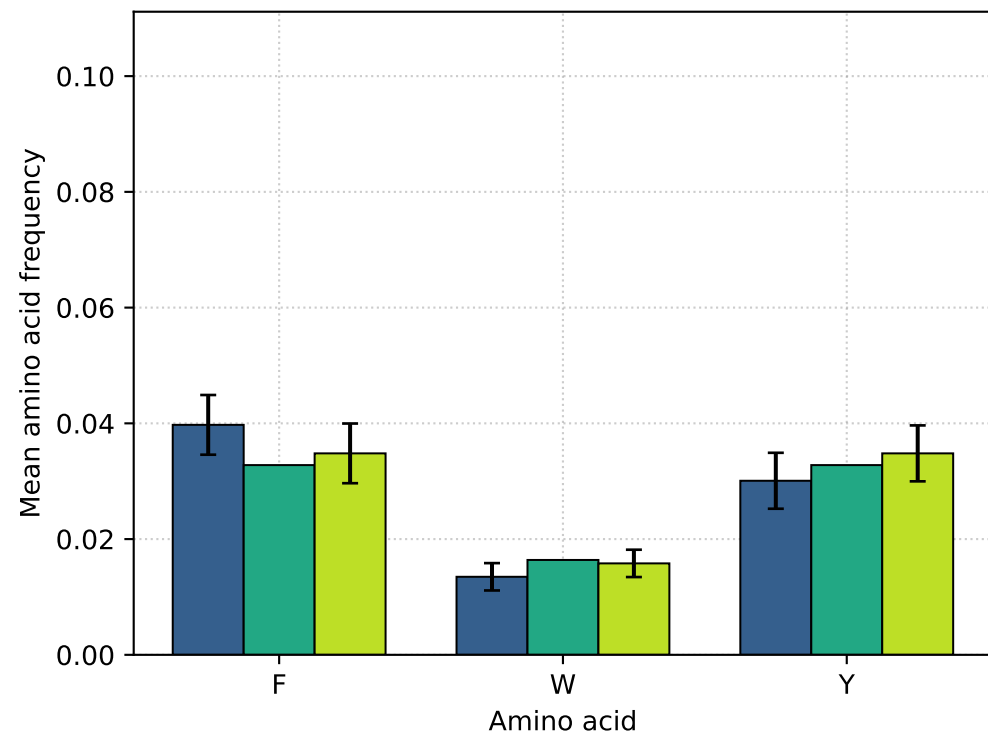
Charged amino acids



Codon + GC correlation:

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

Aromatic amino acids



Uncharged amino acids

