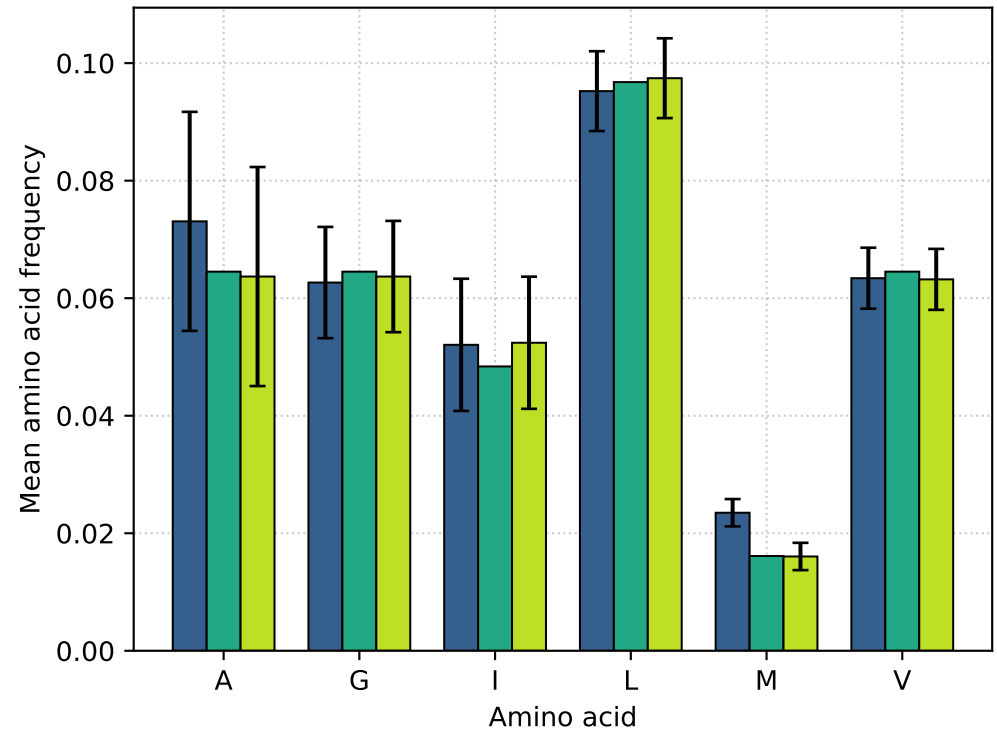


Eukaryota - Mean proteome amino acid frequency for genetic code: Euplotid nuclear

Aliphatic amino acids



Observed
Codon
Codon+GC

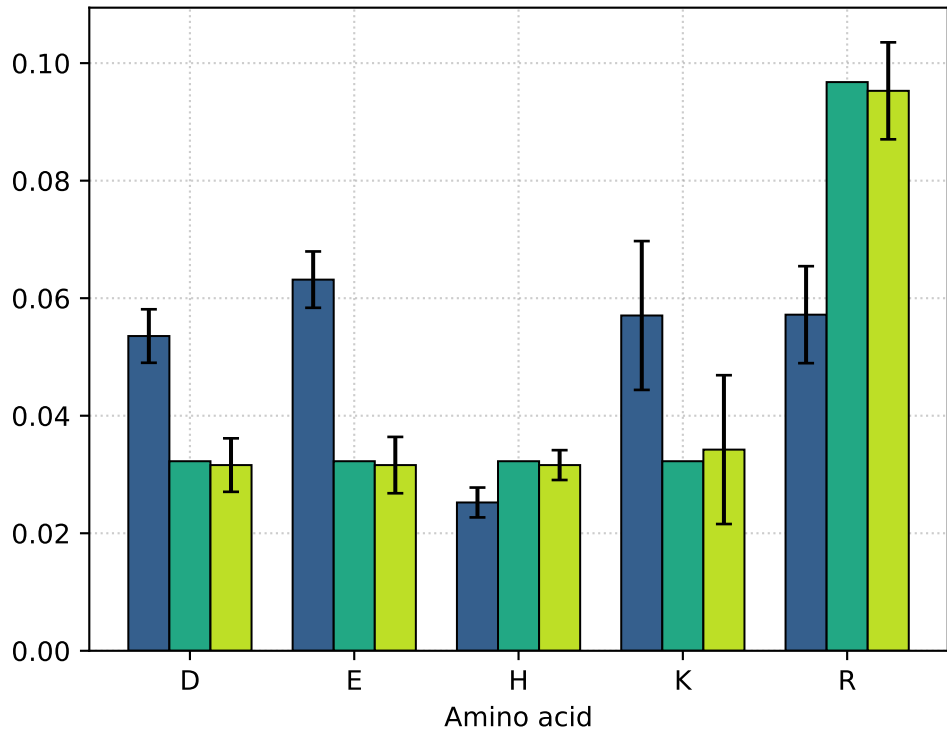
Codon correlation:

- Mean Pearson:
- Coefficient: 0.71552
- p-value: 1.68266e-02

- Mean Spearman:
- Coefficient: 0.69259
- p-value: 1.87519e-02

- Mean log-MSE: 0.33854

Charged amino acids



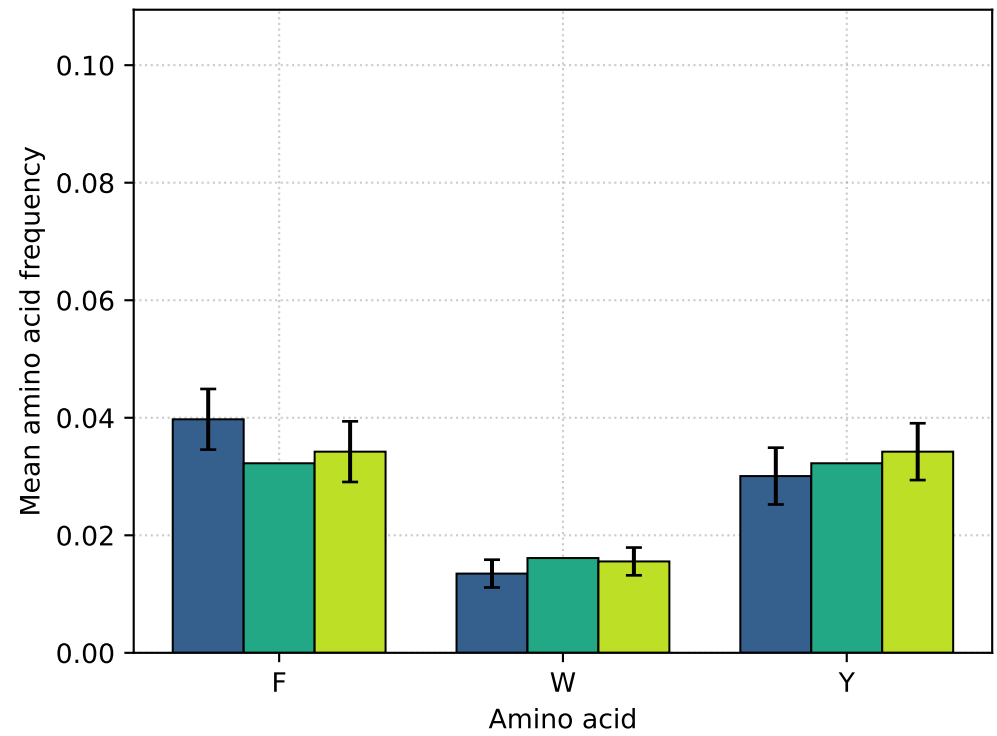
Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.75707
- p-value: 3.57004e-04

- Mean Spearman:
- Coefficient: 0.71064
- p-value: 1.89950e-03

- Mean log-MSE: 0.32462

Aromatic amino acids



Uncharged amino acids

