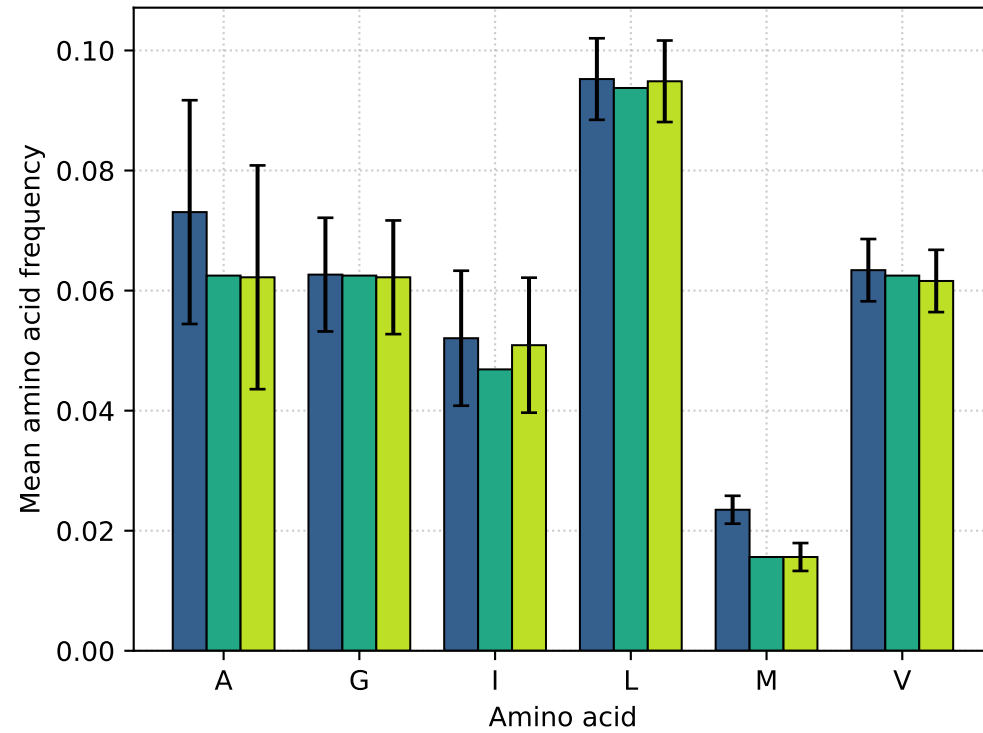


Eukaryota - Mean proteome amino acid frequency for genetic code: Karyorelict nuclear

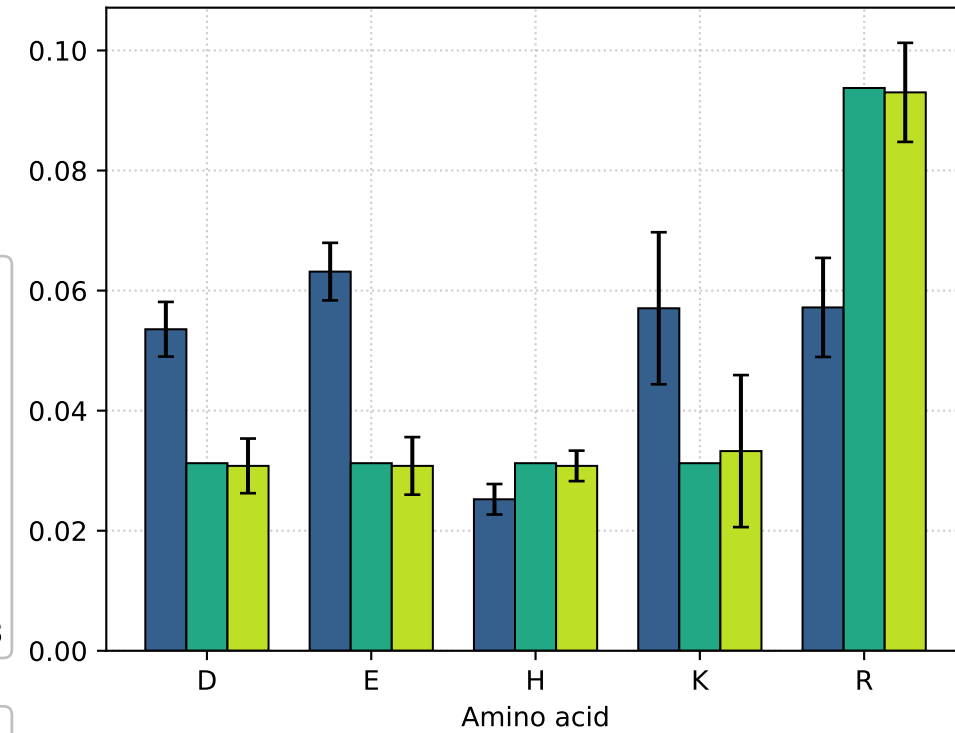
Aliphatic amino acids



Codon correlation:

- Mean Pearson:
- Coefficient: 0.69782
- p-value: 1.68519e-02
- Mean Spearman:
- Coefficient: 0.67630
- p-value: 2.28897e-02
- Mean log-MSE: 0.36868

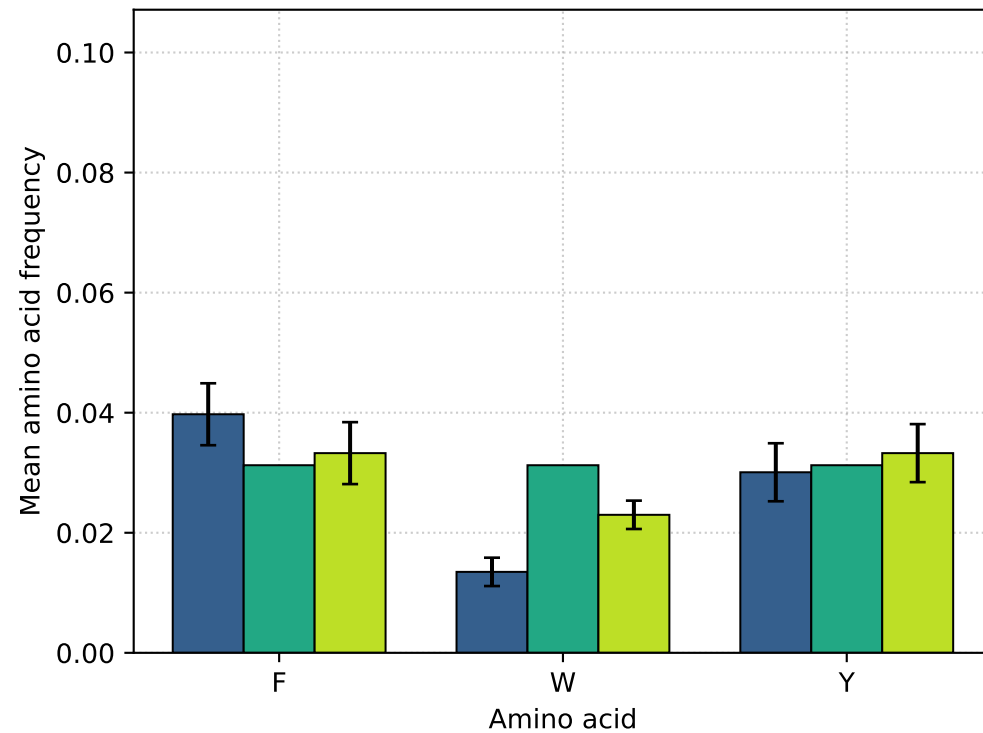
Charged amino acids



Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.75041
- p-value: 5.15894e-04
- Mean Spearman:
- Coefficient: 0.72230
- p-value: 1.79354e-03
- Mean log-MSE: 0.31423

Aromatic amino acids



Uncharged amino acids

