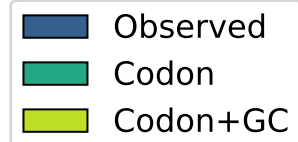
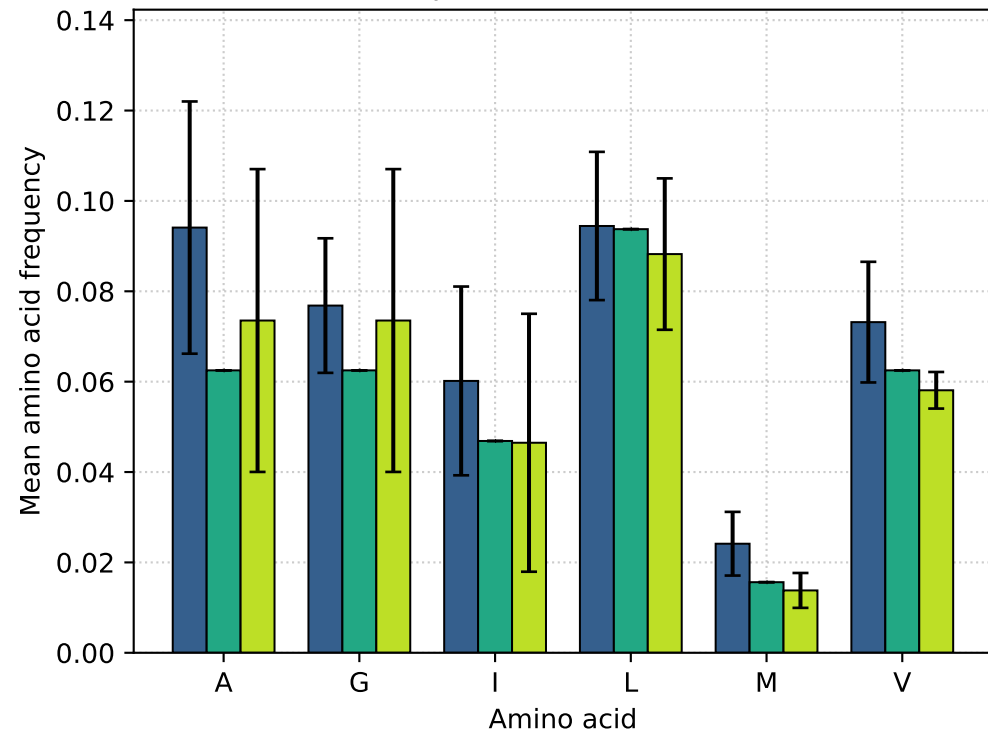


Bacteria - Mean proteomic amino acid frequency for genetic code: Euplotid nuclear

Aliphatic amino acids



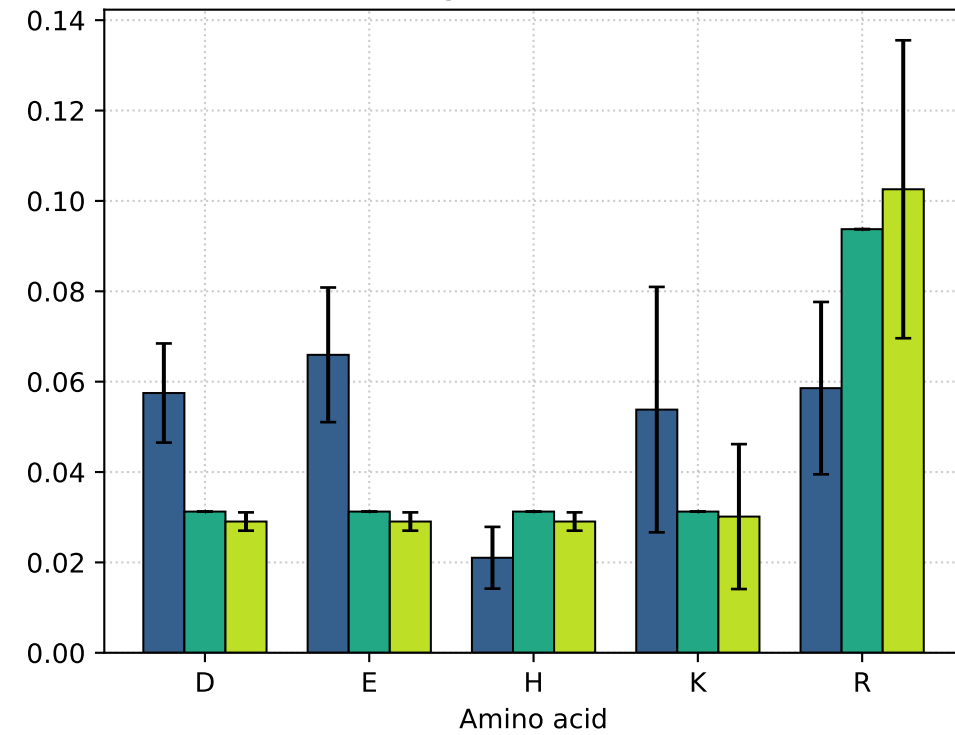
Codon correlation:

- Pearson:
- Coefficient: 0.529429
- p-value: 0.058017
- Spearman:
- Coefficient: 0.563590
- p-value: 0.044143

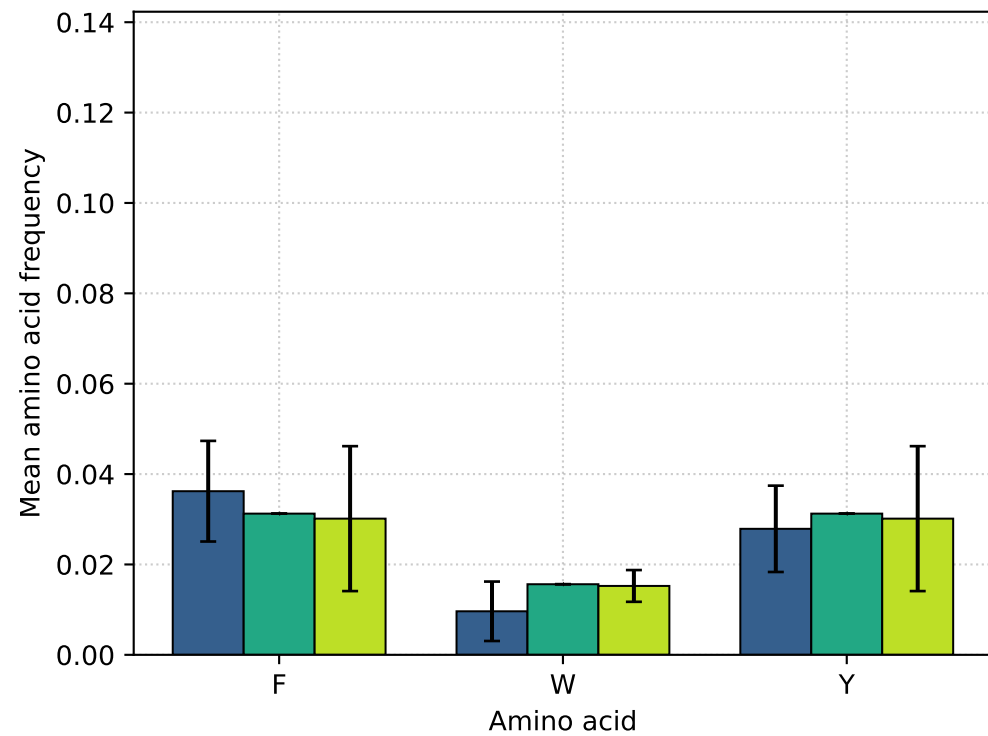
Codon + GC correlation:

- Pearson:
- Coefficient: 0.599261
- p-value: 0.021607
- Spearman:
- Coefficient: 0.586140
- p-value: 0.027802

Charged amino acids



Aromatic amino acids



Uncharged amino acids

