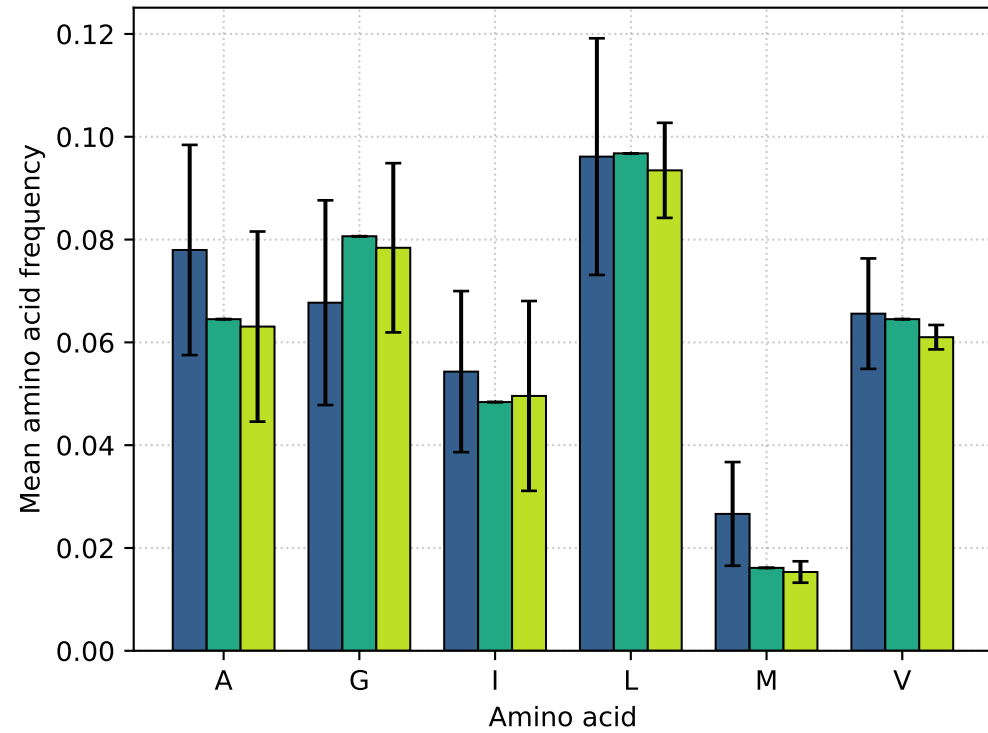


Eukaryota - Mean proteome amino acid frequency for genetic code: Candidate division sr1 and gracilibacteria

Aliphatic amino acids



Observed
Codon
Codon+GC

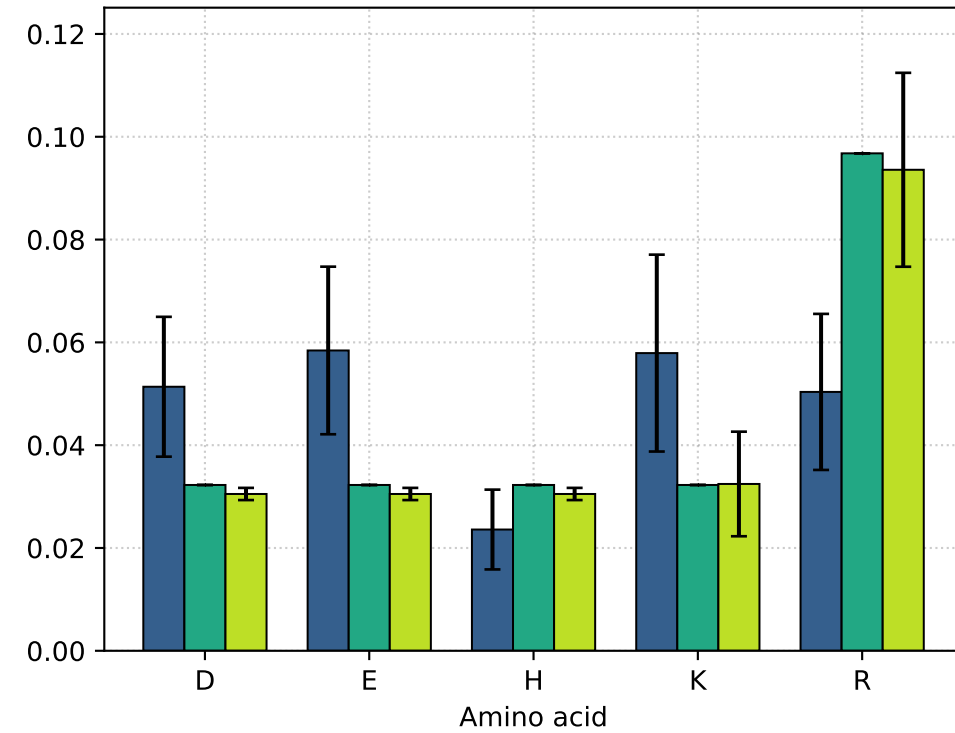
Codon correlation:
- Mean Pearson:
- Coefficient: 0.64824
- p-value: 2.24232e-02

- Mean Spearman:
- Coefficient: 0.67069
- p-value: 1.52262e-02

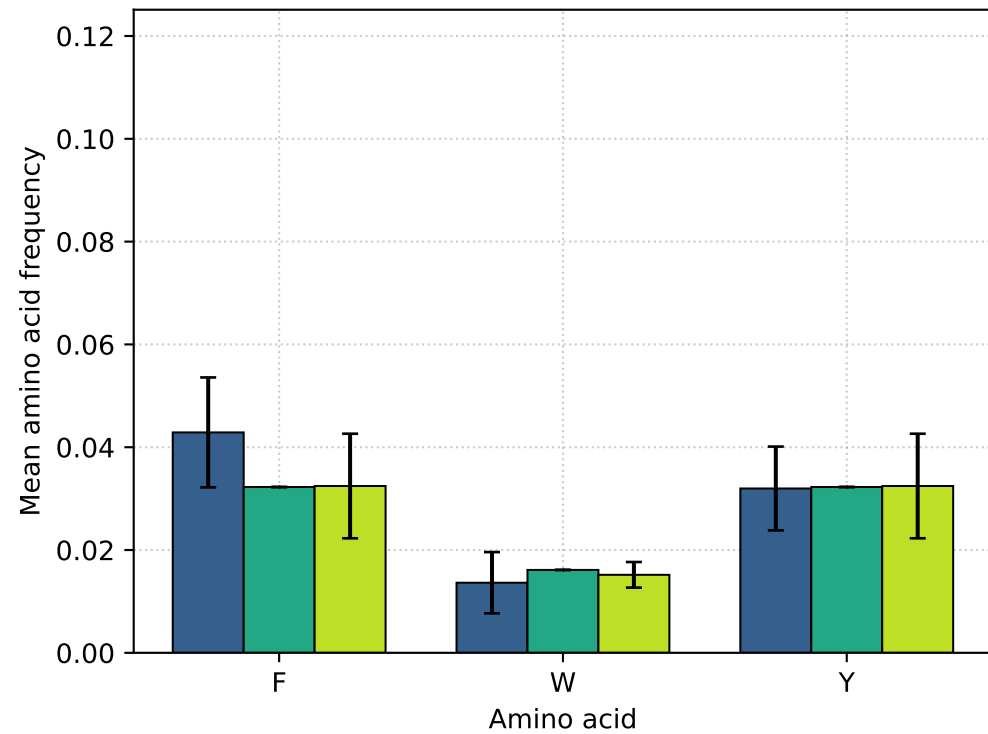
Codon + GC correlation:
- Mean Pearson:
- Coefficient: 0.67680
- p-value: 1.06482e-02

- Mean Spearman:
- Coefficient: 0.67145
- p-value: 9.24144e-03

Charged amino acids



Aromatic amino acids



Uncharged amino acids

