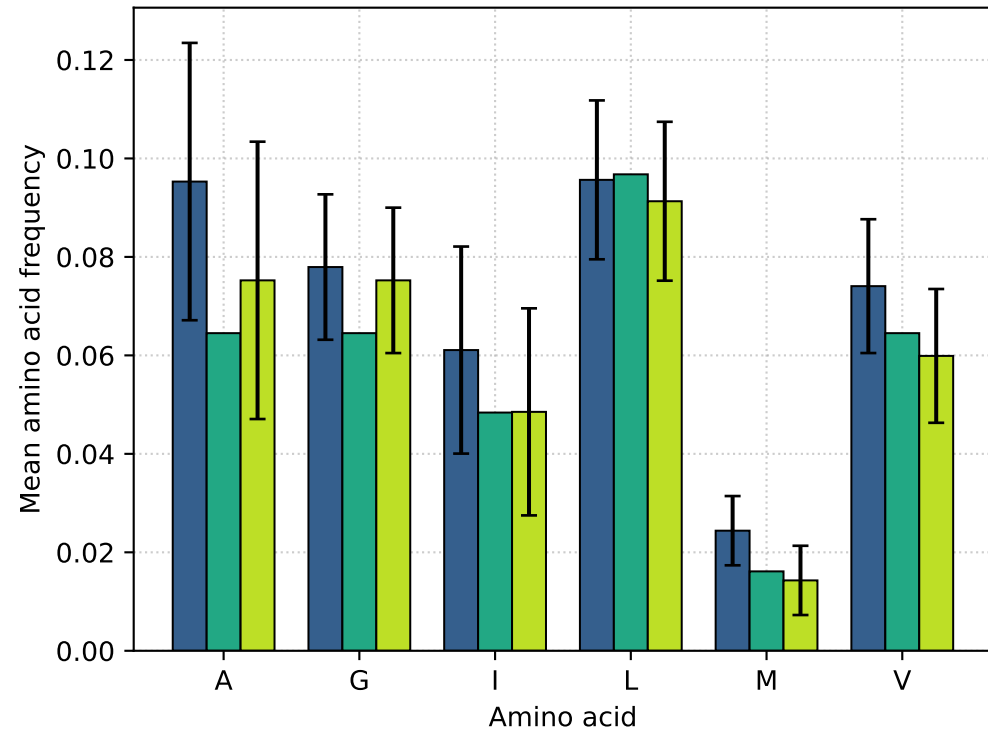


Bacteria - Mean proteome amino acid frequency for genetic code: Ciliate-dasycladacean-hexamita nuclear

Aliphatic amino acids

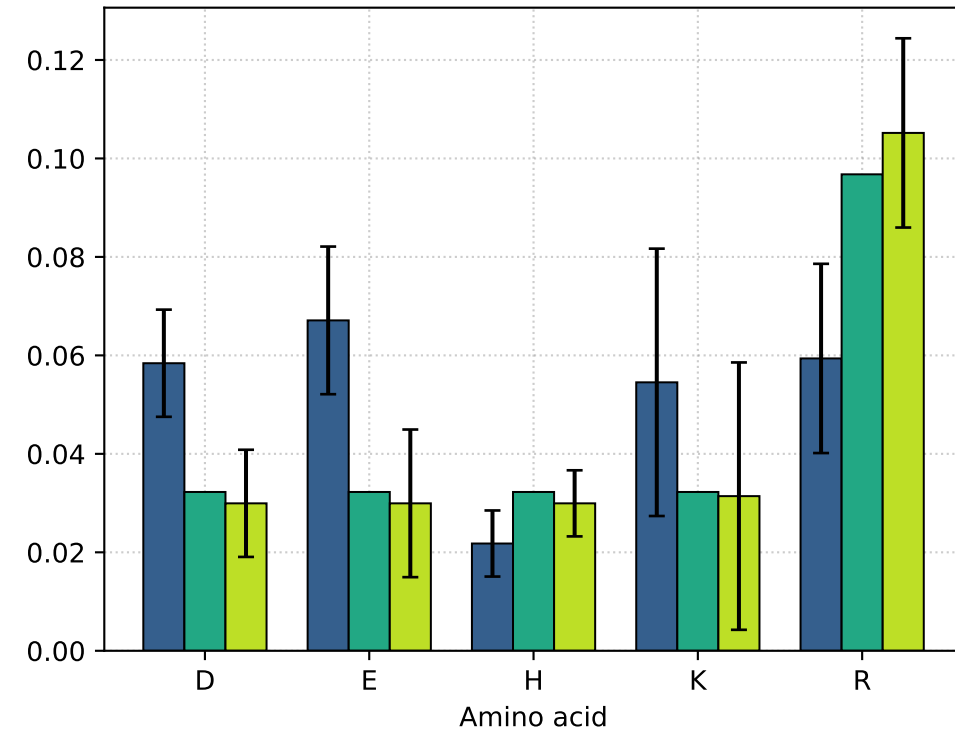


Observed
Codon
Codon+GC

Codon correlation:
- Mean Pearson:
- Coefficient: 0.55809
- p-value: 4.80889e-02

- Mean Spearman:
- Coefficient: 0.60845
- p-value: 3.27802e-02

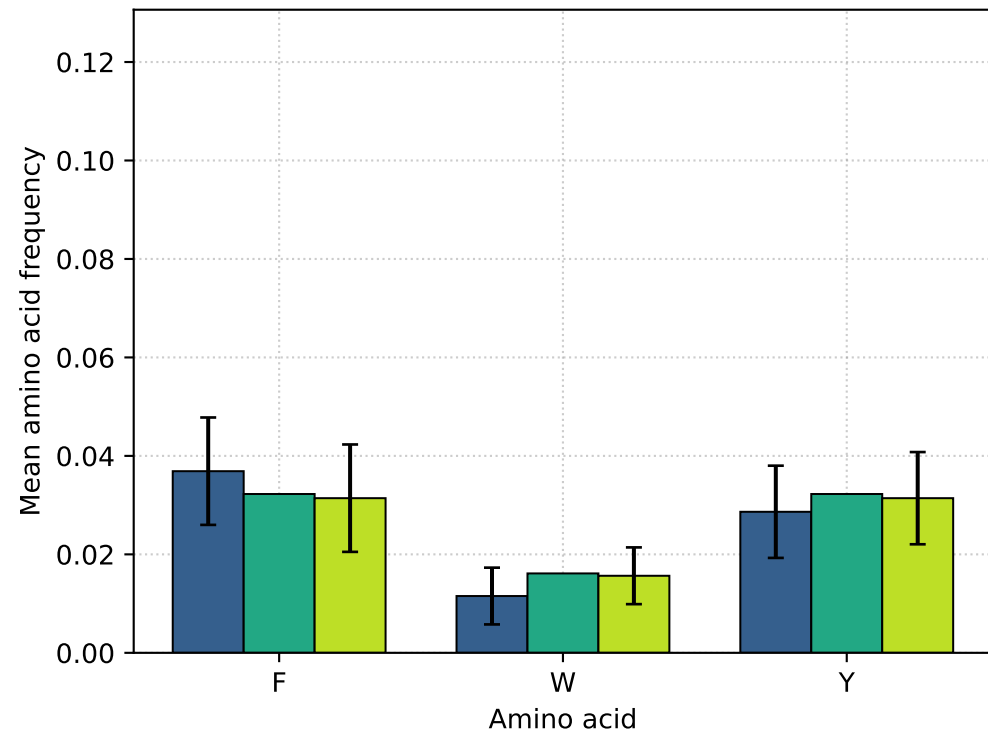
Charged amino acids



Codon + GC correlation:
- Mean Pearson:
- Coefficient: 0.62434
- p-value: 1.67772e-02

- Mean Spearman:
- Coefficient: 0.62363
- p-value: 1.78423e-02

Aromatic amino acids



Uncharged amino acids

