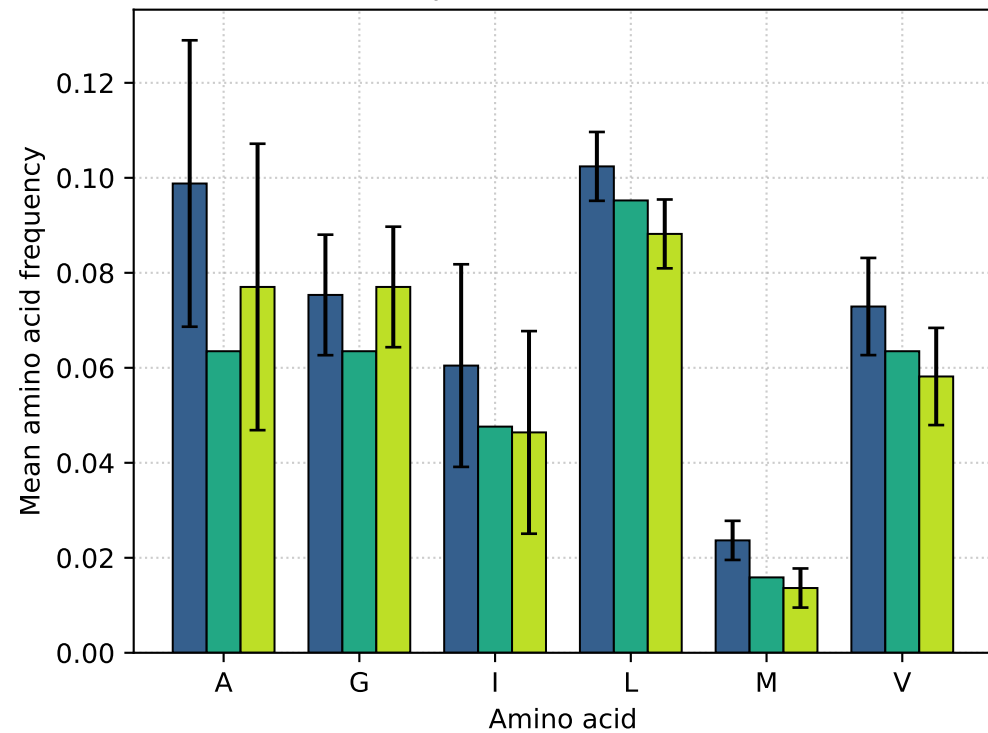
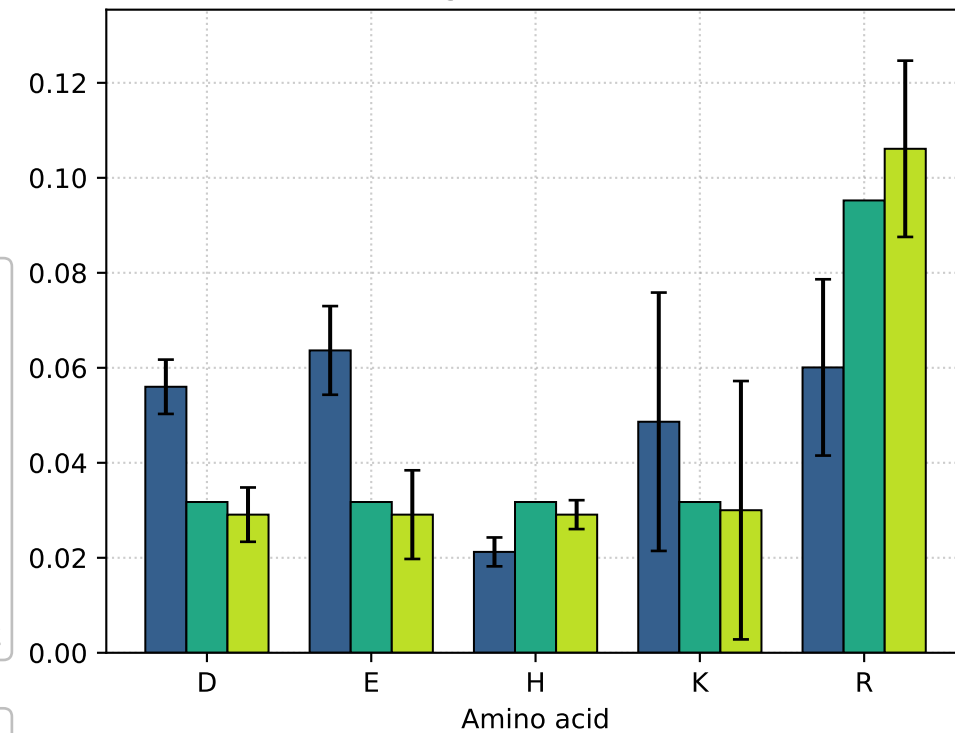


Bacteria - Mean proteome amino acid frequency for genetic code: Mesodinium nuclear

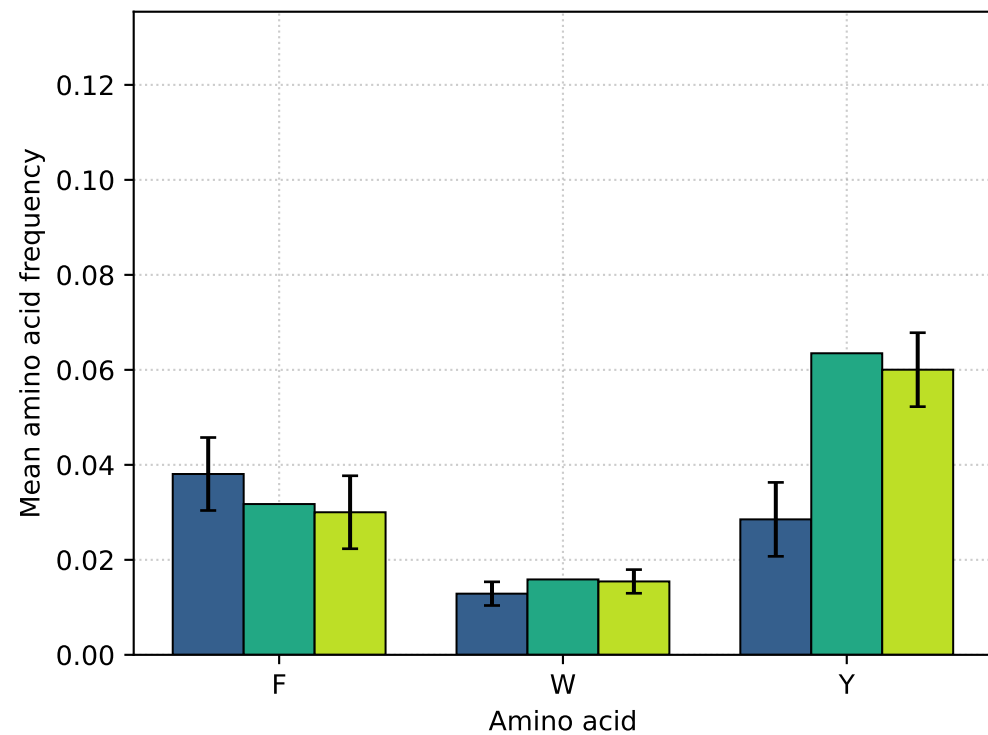
Aliphatic amino acids



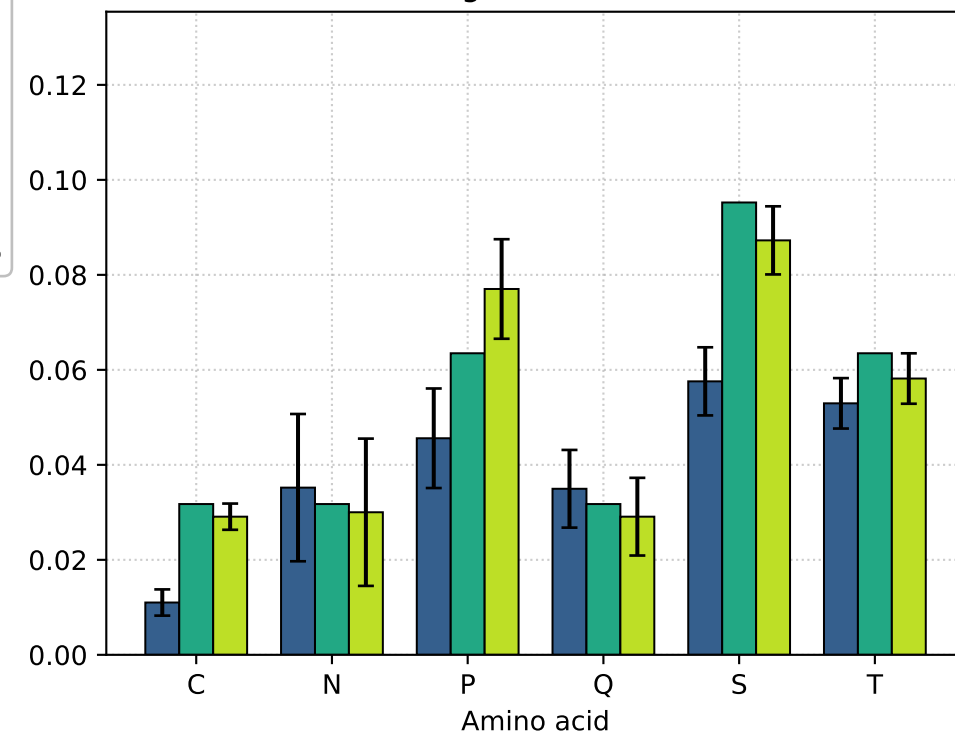
Charged amino acids



Aromatic amino acids



Uncharged amino acids



Codon correlation:

- Mean Pearson:
- Coefficient: 0.57641
- p-value: 3.31414e-02
- Mean Spearman:
- Coefficient: 0.60003
- p-value: 2.63460e-02
- Mean log-MSE: 0.56662

Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.65717
- p-value: 4.54149e-03
- Mean Spearman:
- Coefficient: 0.65236
- p-value: 6.02739e-03
- Mean log-MSE: 0.59875