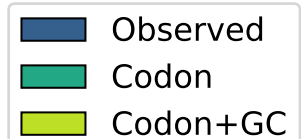
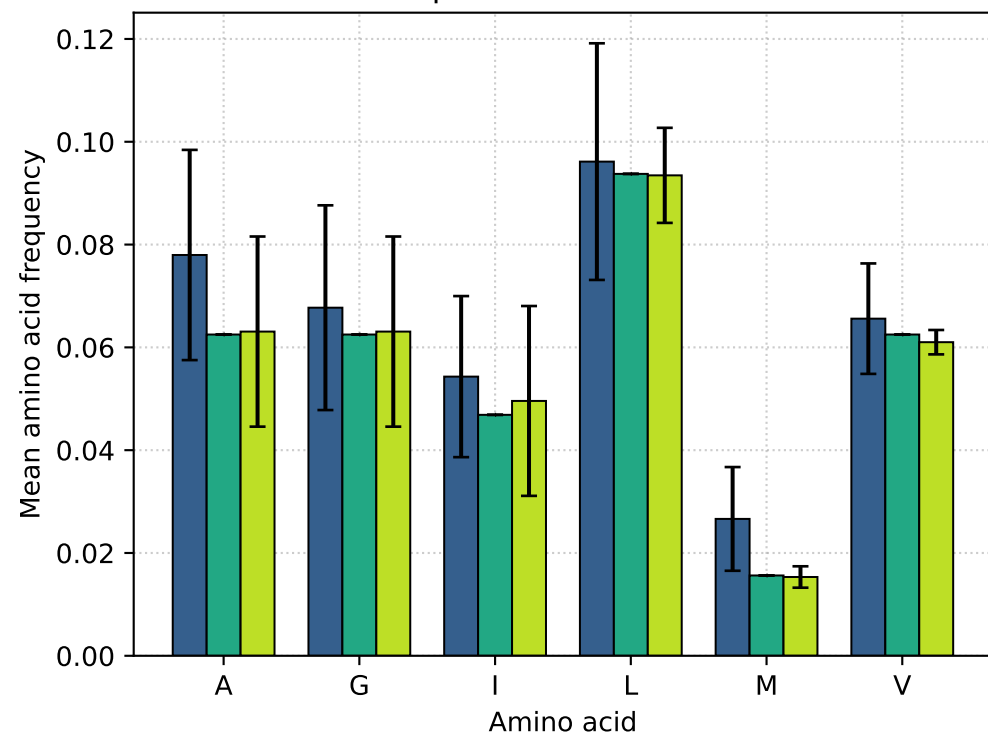


Eukaryota - Mean proteome amino acid frequency for genetic code: Condyllostoma nuclear

Aliphatic amino acids



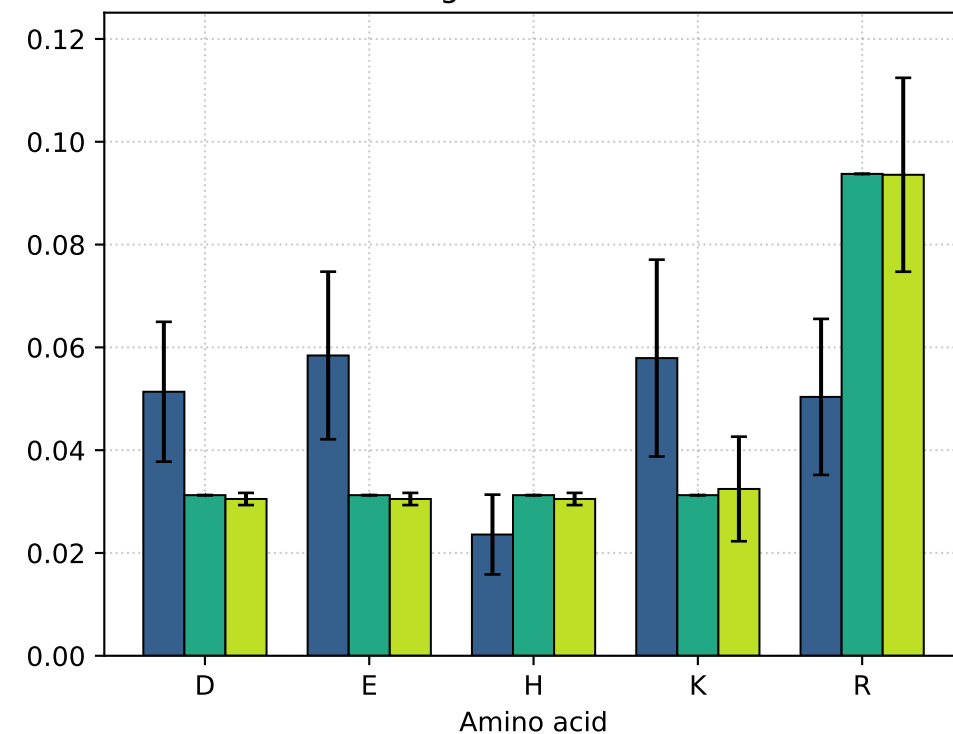
Codon correlation:
- Mean Pearson:
- Coefficient: 0.58742
- p-value: 4.22175e-02

- Mean Spearman:
- Coefficient: 0.58769
- p-value: 4.05337e-02

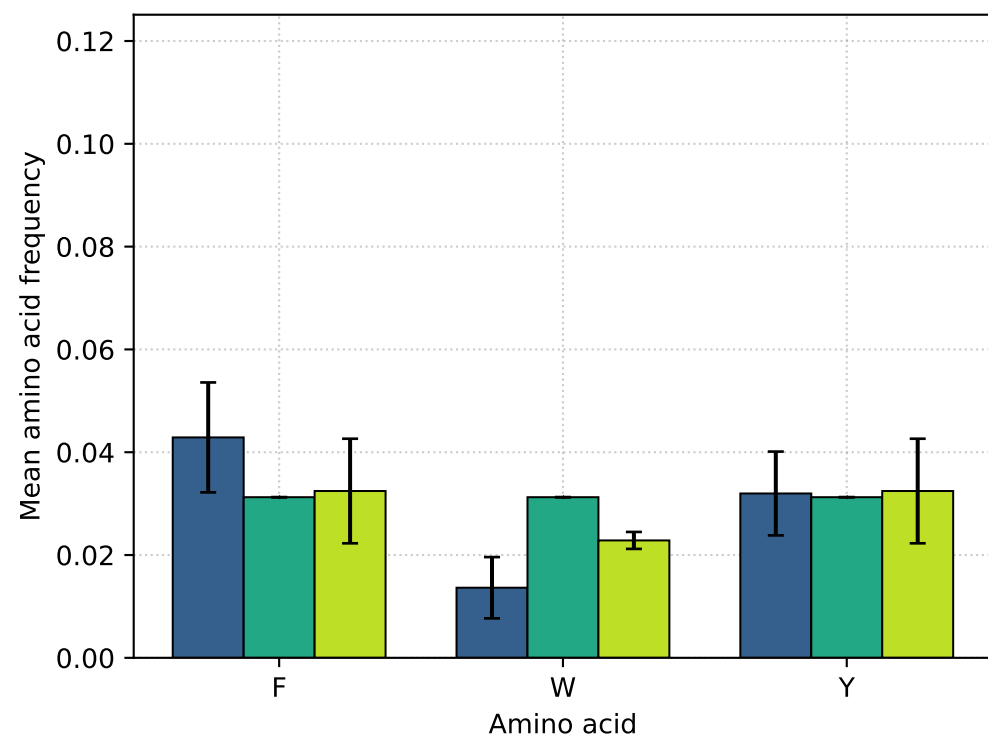
Codon + GC correlation:
- Mean Pearson:
- Coefficient: 0.64970
- p-value: 1.54782e-02

- Mean Spearman:
- Coefficient: 0.63950
- p-value: 1.50610e-02

Charged amino acids



Aromatic amino acids



Uncharged amino acids

