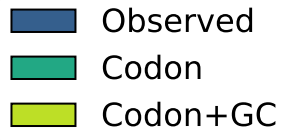
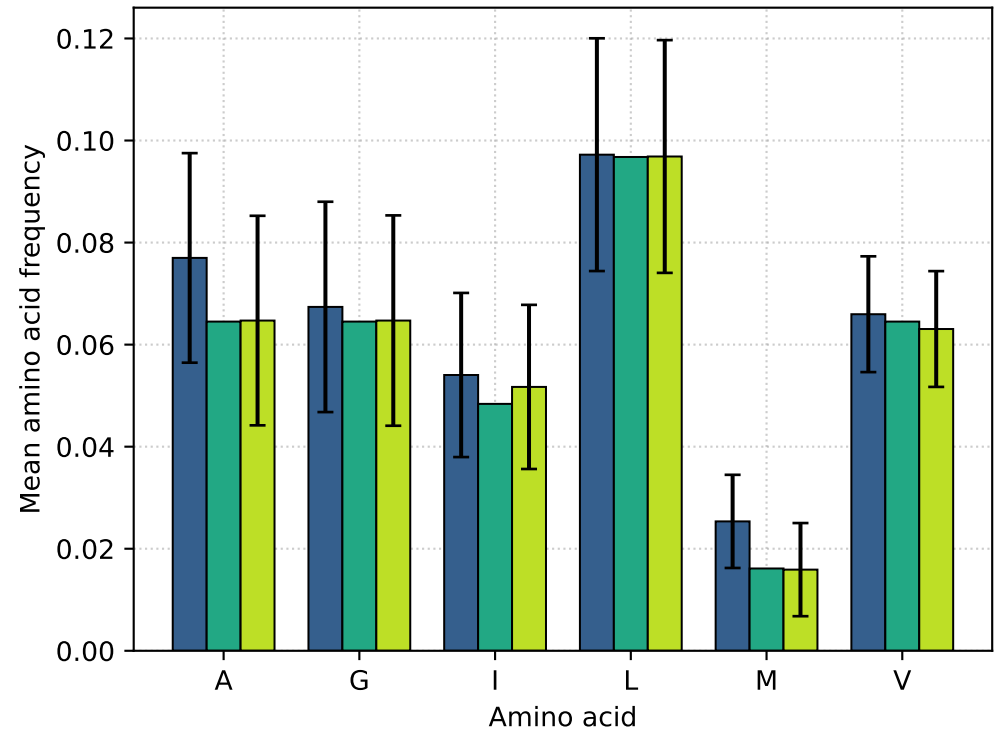


# Eukaryota - Mean proteome amino acid frequency for genetic code: Euplotid nuclear

Aliphatic amino acids



*Codon correlation:*

- Mean Pearson:
- Coefficient: 0.60350
- p-value: 3.12328e-02

- Mean Spearman:

- Coefficient: 0.60446
- p-value: 3.04261e-02

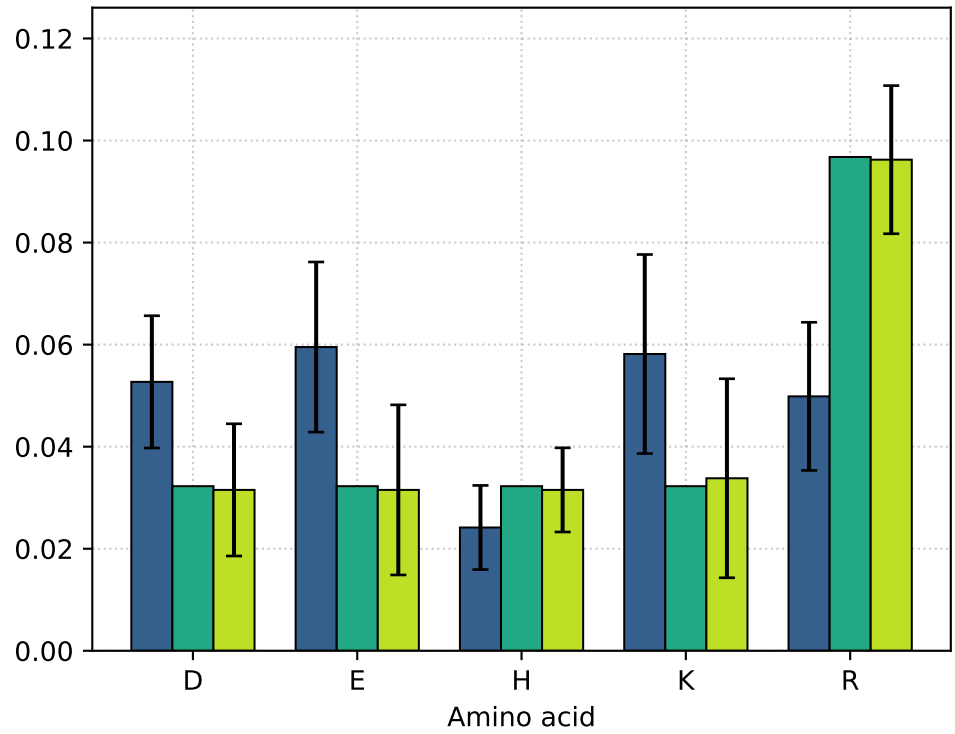
*Codon + GC correlation:*

- Mean Pearson:
- Coefficient: 0.63334
- p-value: 1.74554e-02

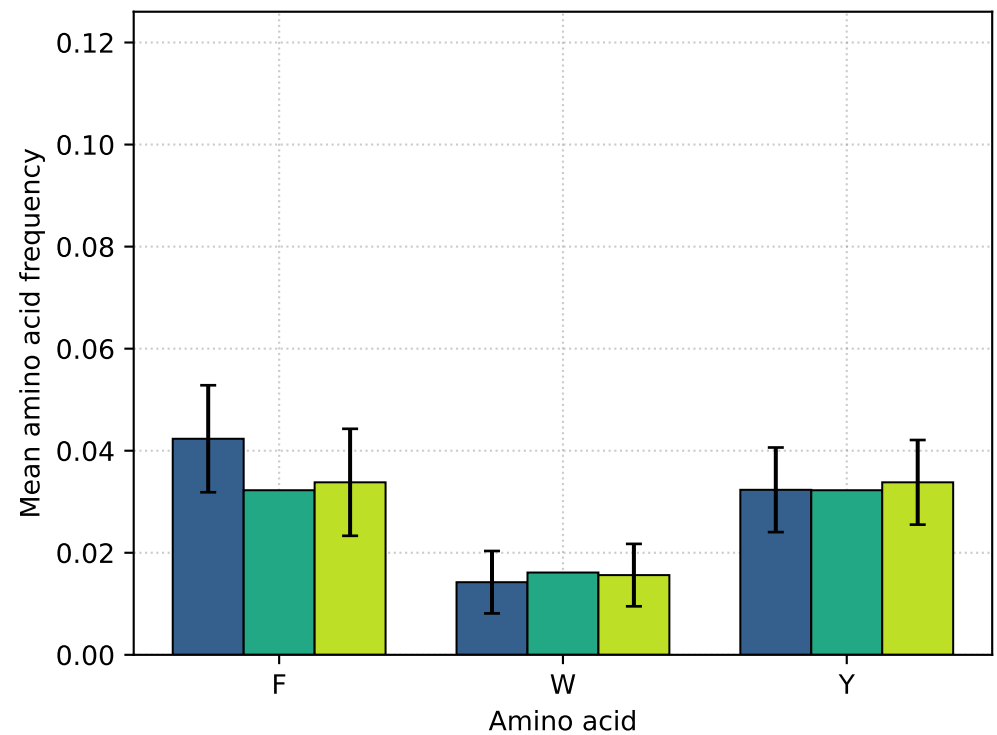
- Mean Spearman:

- Coefficient: 0.60886
- p-value: 2.08777e-02

Charged amino acids



Aromatic amino acids



Uncharged amino acids

