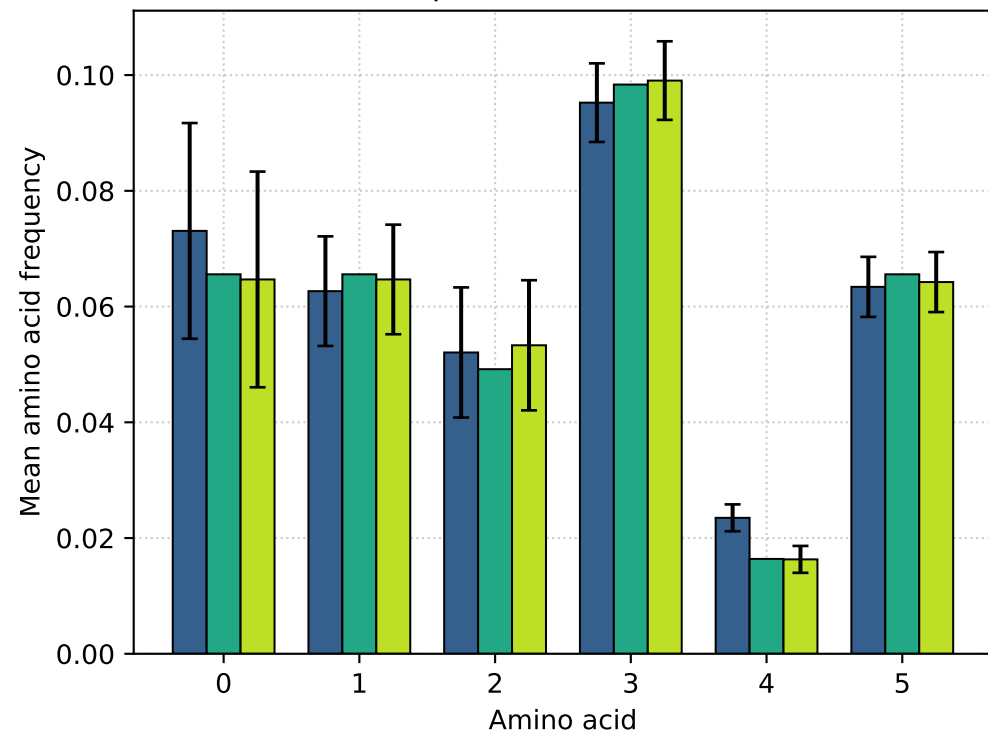


Eukaryota - Mean proteome amino acid frequency for genetic code: Standard

Aliphatic amino acids



Observed
Codon
Codon+GC

Codon correlation:

- Mean Pearson:
- Coefficient: 0.75245
- p-value: 1.38887e-02

- Mean Spearman:
- Coefficient: 0.75366
- p-value: 1.05887e-02

- Mean log-MSE: 0.27275

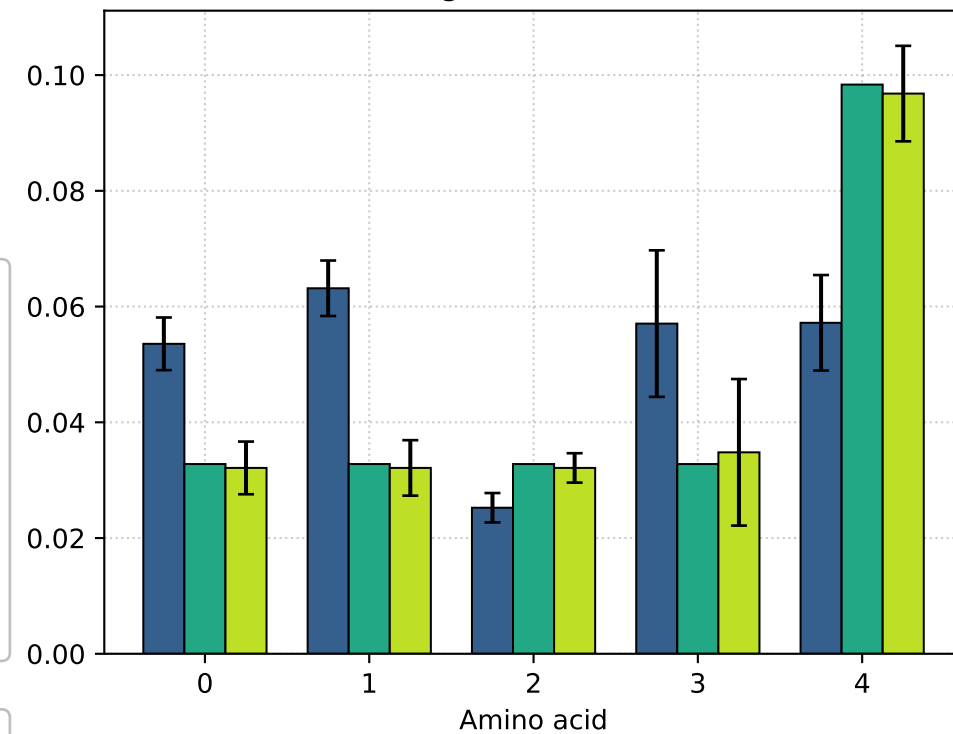
Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.79252
- p-value: 1.62267e-04

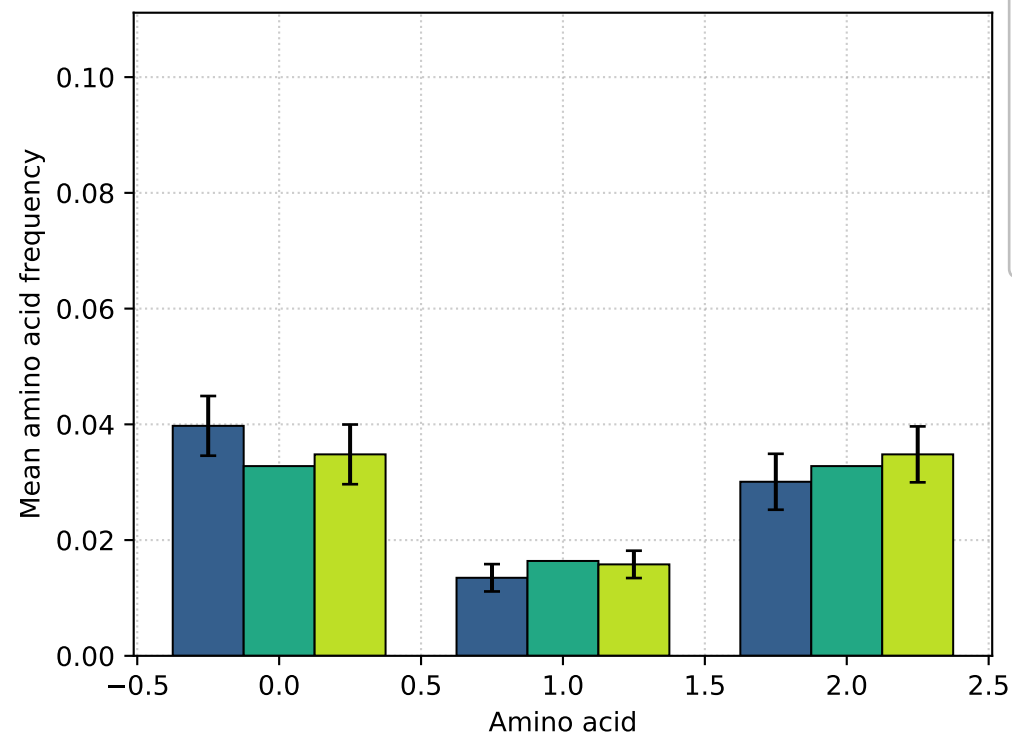
- Mean Spearman:
- Coefficient: 0.76515
- p-value: 3.36228e-04

- Mean log-MSE: 0.25888

Charged amino acids



Aromatic amino acids



Uncharged amino acids

