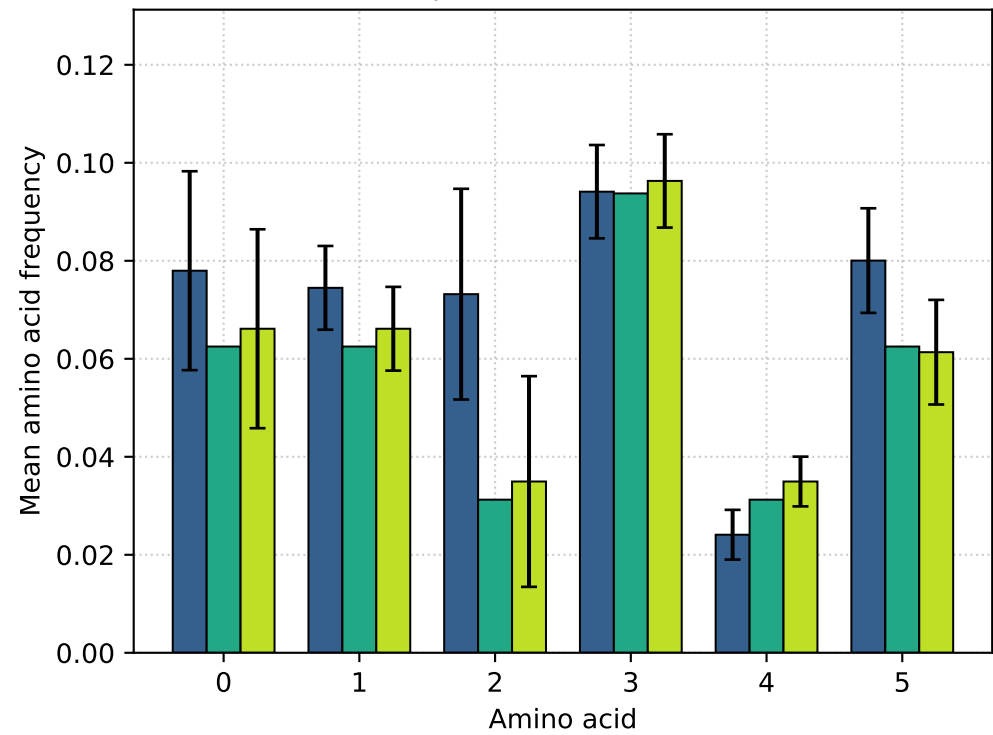
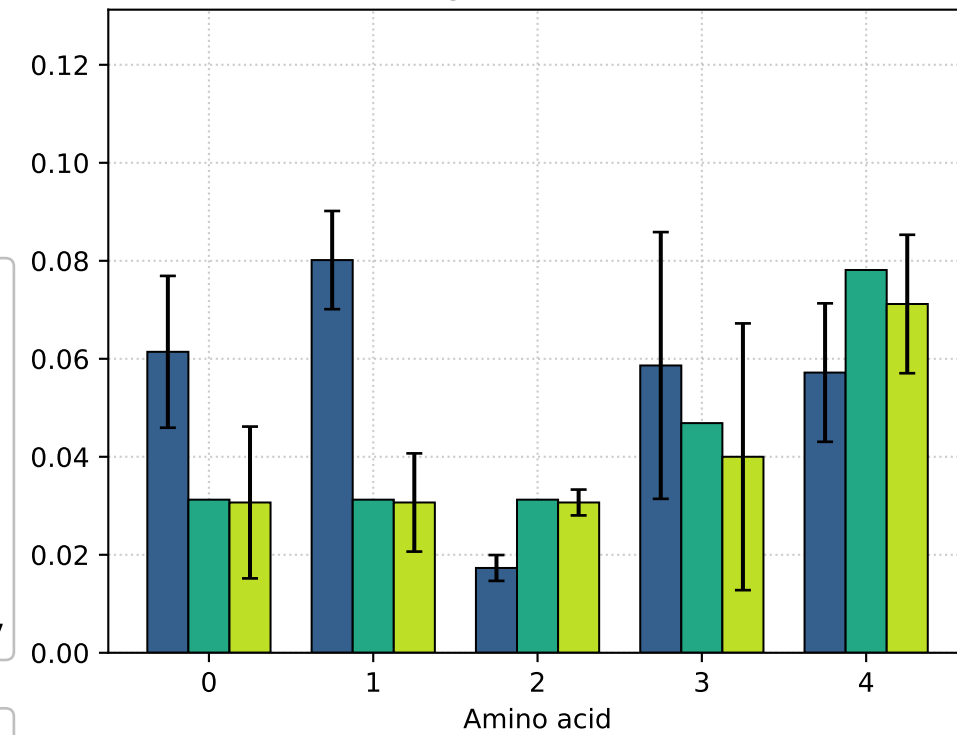


Archaea - Mean proteome amino acid frequency for genetic code: Invertebrate mitochondrial

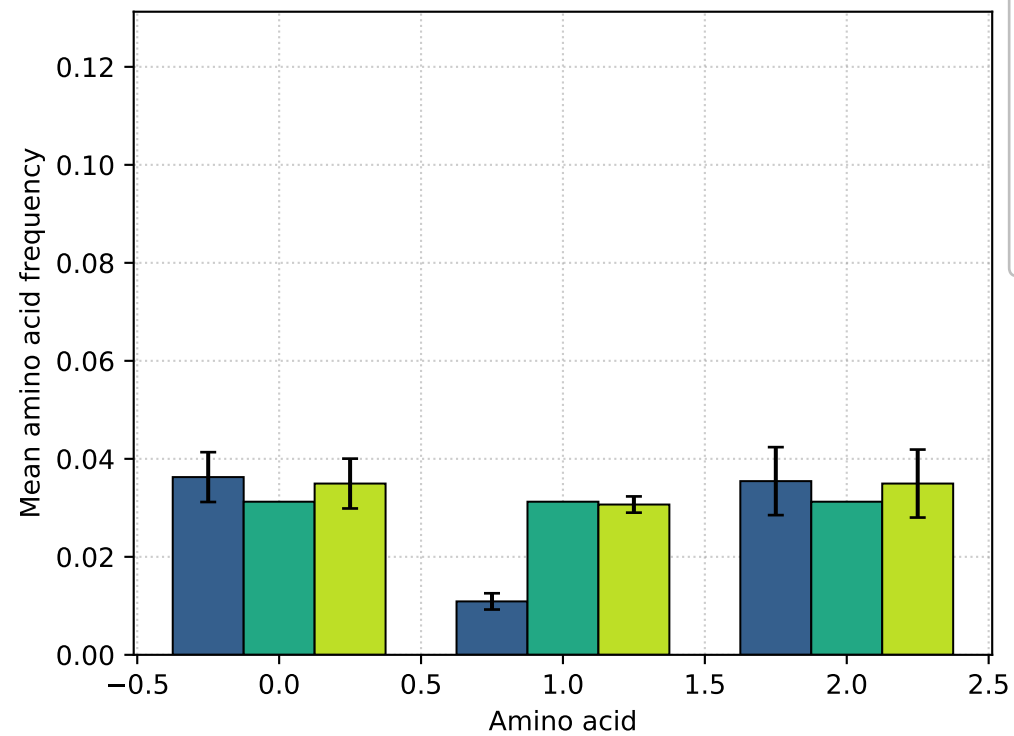
Aliphatic amino acids



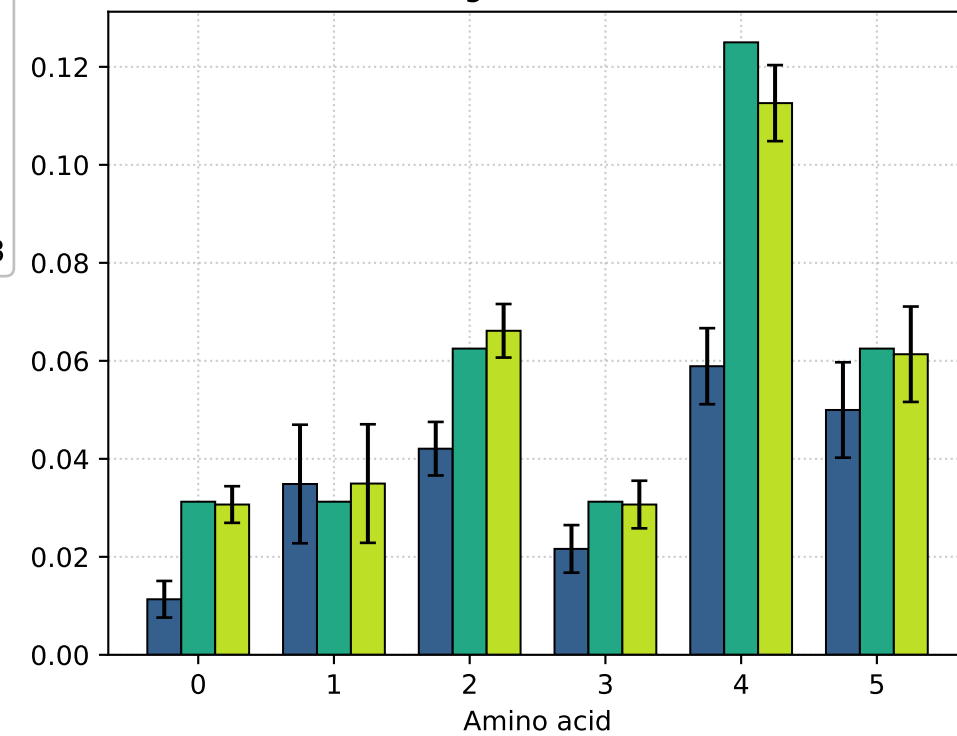
Charged amino acids



Aromatic amino acids



Uncharged amino acids



Codon correlation:
- Mean Pearson:
- Coefficient: 0.45900
- p-value: 7.41339e-02

- Mean Spearman:
- Coefficient: 0.53129
- p-value: 3.85711e-02
- Mean log-MSE: 0.73097

Codon + GC correlation:
- Mean Pearson:
- Coefficient: 0.53831
- p-value: 2.43182e-02

- Mean Spearman:
- Coefficient: 0.52795
- p-value: 2.56899e-02
- Mean log-MSE: 0.73898