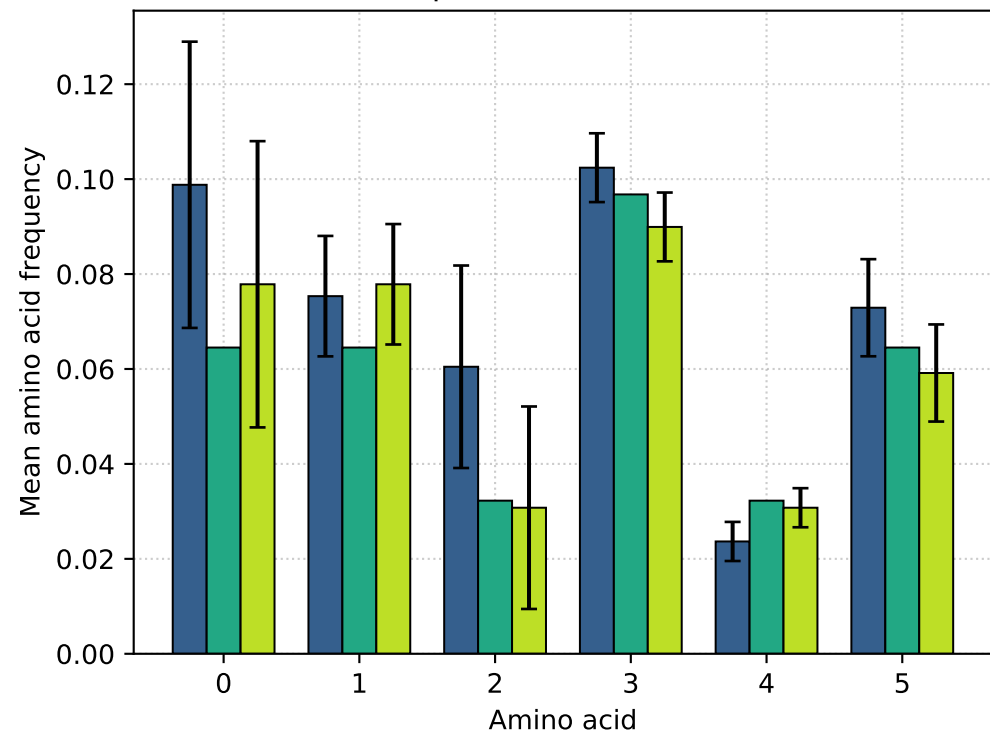


Bacteria - Mean proteome amino acid frequency for genetic code: Trematode mitochondrial

Aliphatic amino acids



Observed

Codon

Codon+GC

Codon correlation:

- Mean Pearson:

- Coefficient: 0.49338

- p-value: 7.06494e-02

- Mean Spearman:

- Coefficient: 0.53251

- p-value: 8.75522e-02

- Mean log-MSE: 0.72701

Codon + GC correlation:

- Mean Pearson:

- Coefficient: 0.58383

- p-value: 2.95220e-02

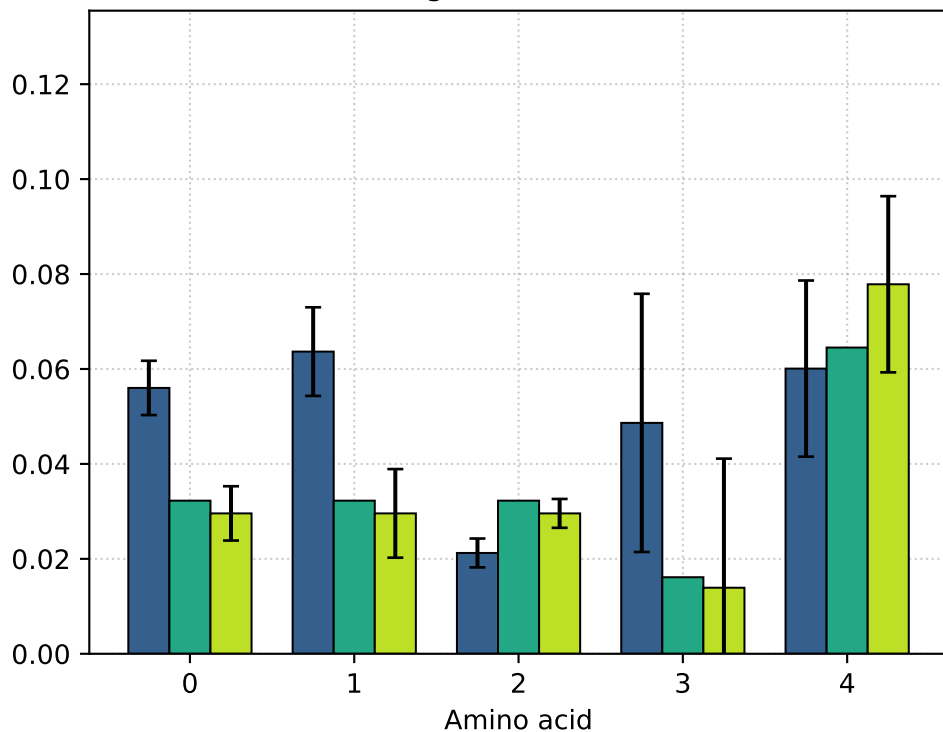
- Mean Spearman:

- Coefficient: 0.53430

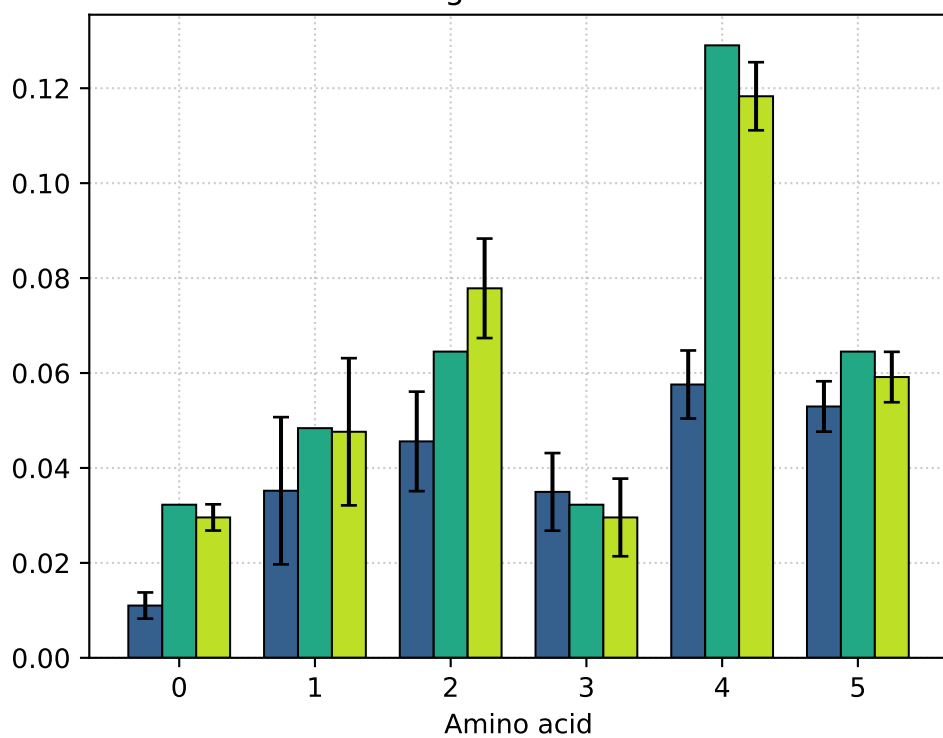
- p-value: 6.93315e-02

- Mean log-MSE: 0.79506

Charged amino acids



Uncharged amino acids



Aromatic amino acids

