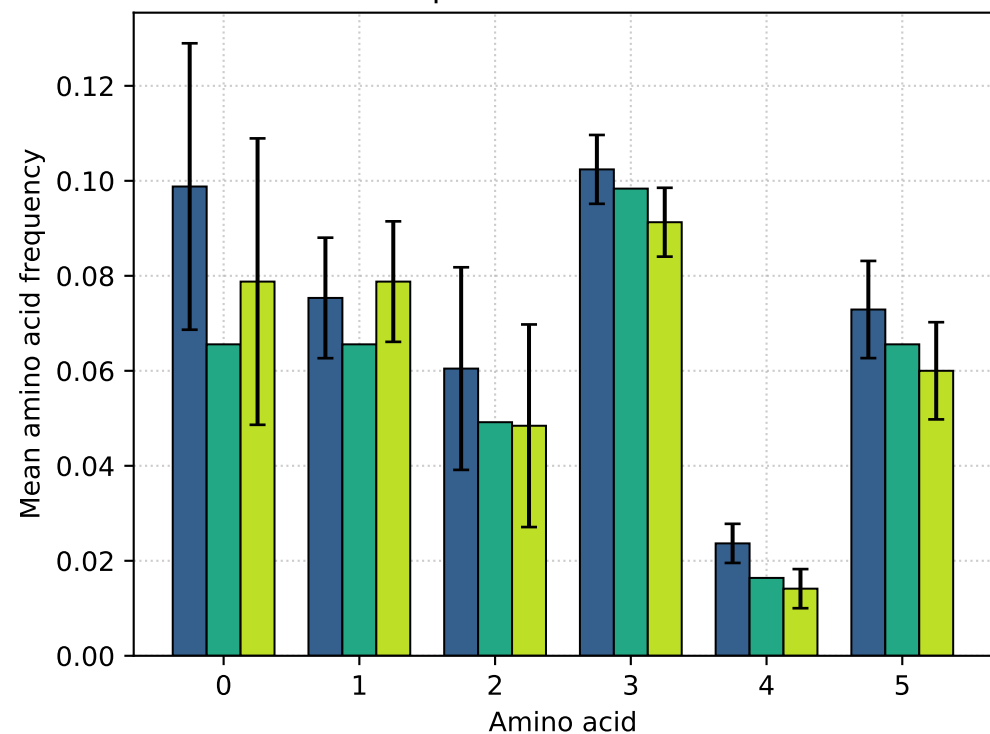
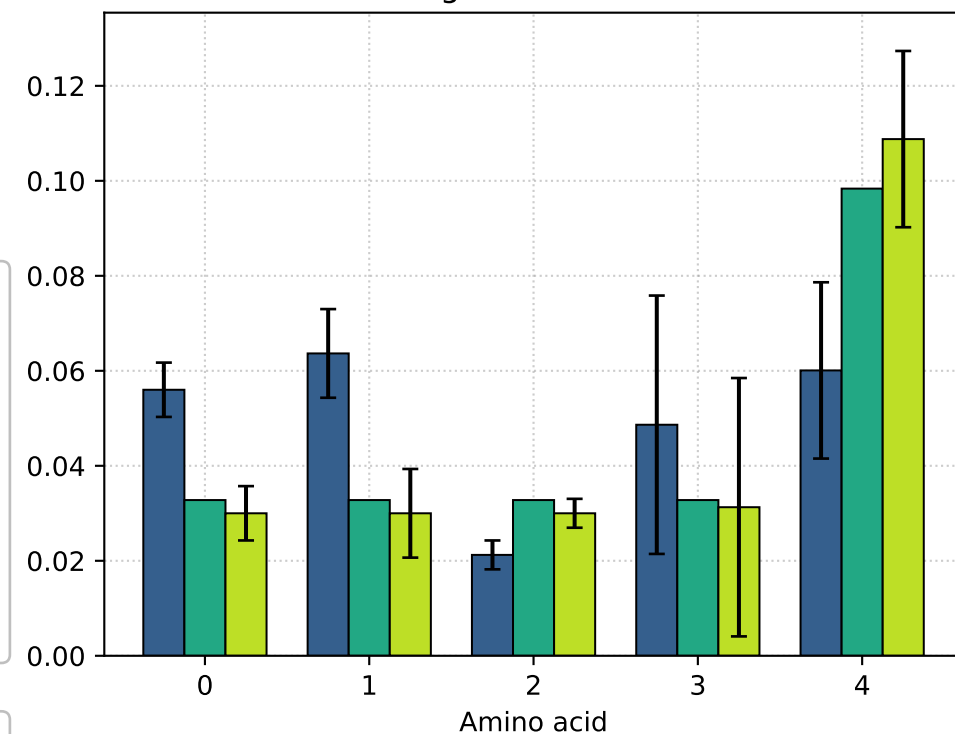


Bacteria - Mean proteome amino acid frequency for genetic code: Bacterial-archaeal-plant plastid

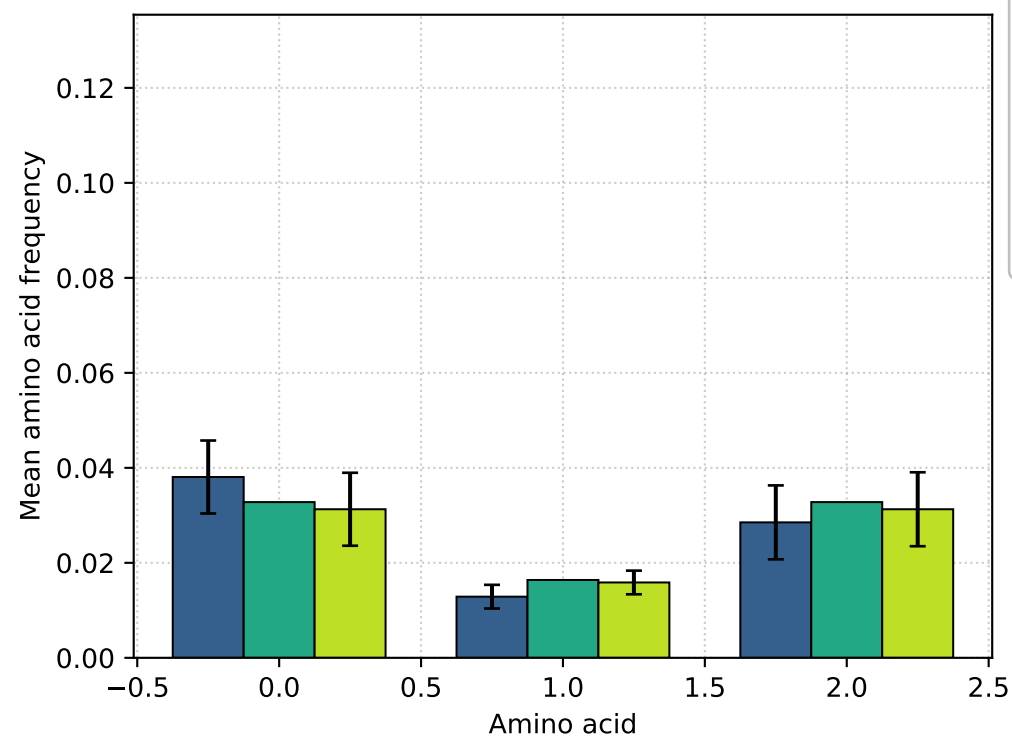
Aliphatic amino acids



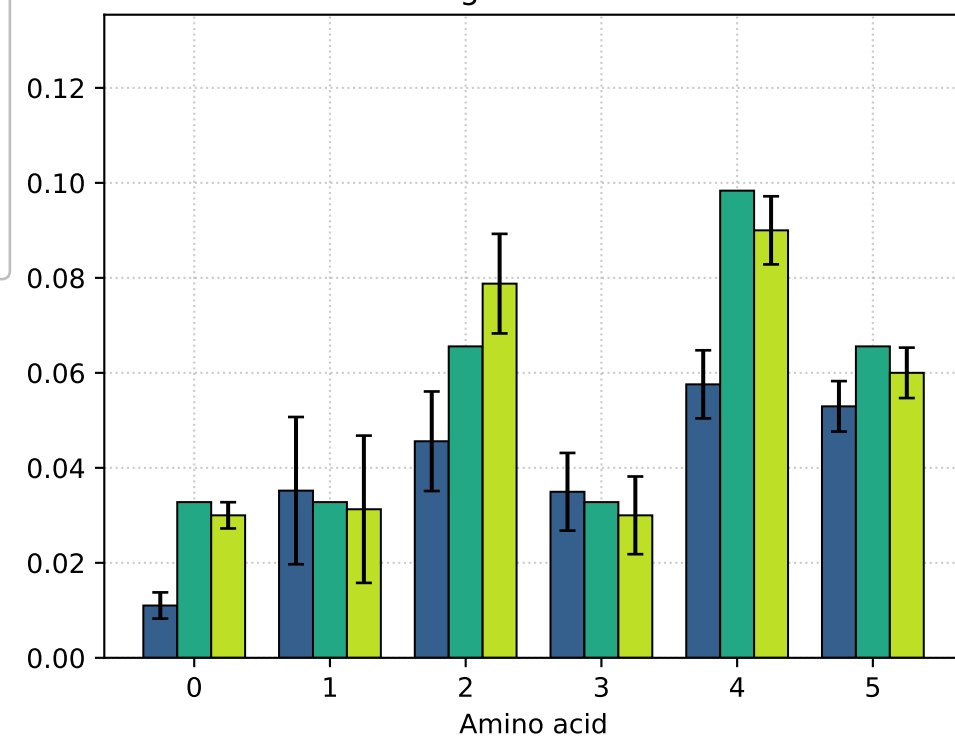
Charged amino acids



Aromatic amino acids



Uncharged amino acids



Codon correlation:

- Mean Pearson:
- Coefficient: 0.62296
- p-value: 2.62645e-02
- Mean Spearman:
- Coefficient: 0.67283
- p-value: 1.56294e-02
- Mean log-MSE: 0.50066

Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.70792
- p-value: 1.11376e-03
- Mean Spearman:
- Coefficient: 0.70086
- p-value: 2.00455e-03
- Mean log-MSE: 0.55671