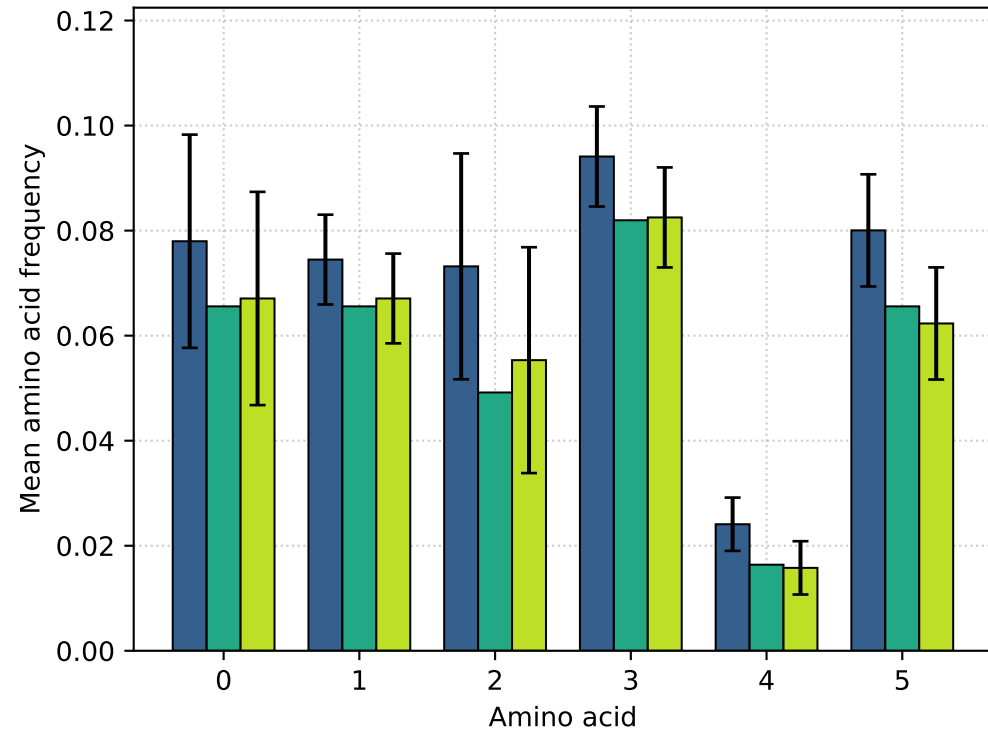
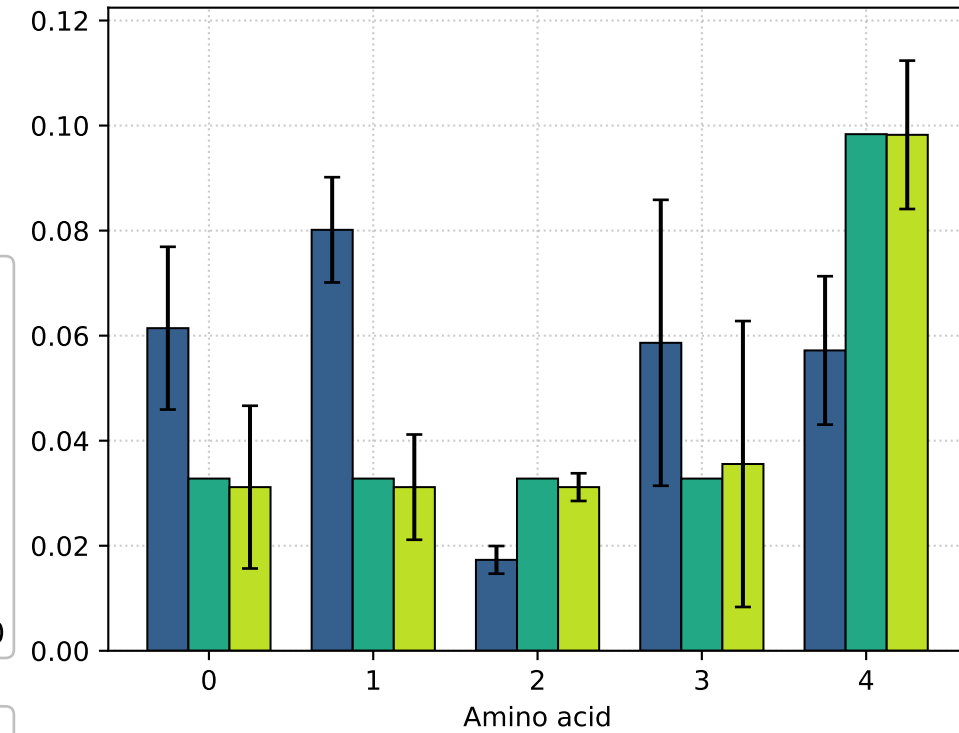


Archaea - Mean proteome amino acid frequency for genetic code: Alternative yeast

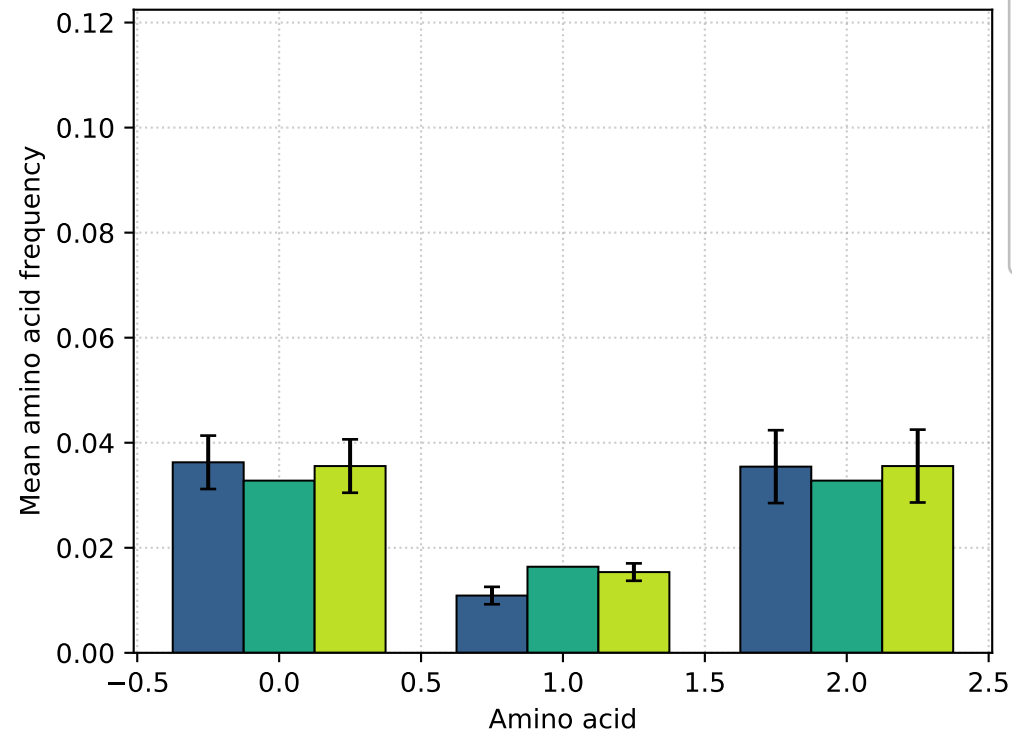
Aliphatic amino acids



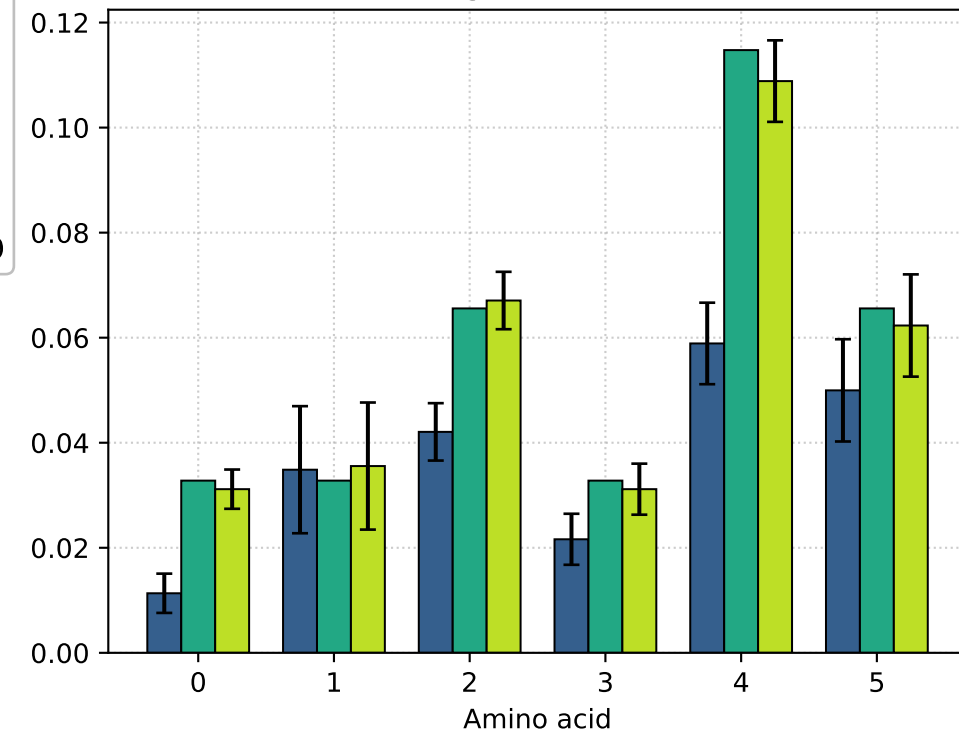
Charged amino acids



Aromatic amino acids



Uncharged amino acids



Codon correlation:

- Mean Pearson:
- Coefficient: 0.50674
- p-value: 5.61321e-02
- Mean Spearman:
- Coefficient: 0.60377
- p-value: 1.75392e-02
- Mean log-MSE: 0.61800

Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.59500
- p-value: 9.25052e-03
- Mean Spearman:
- Coefficient: 0.62889
- p-value: 4.59777e-03
- Mean log-MSE: 0.62009