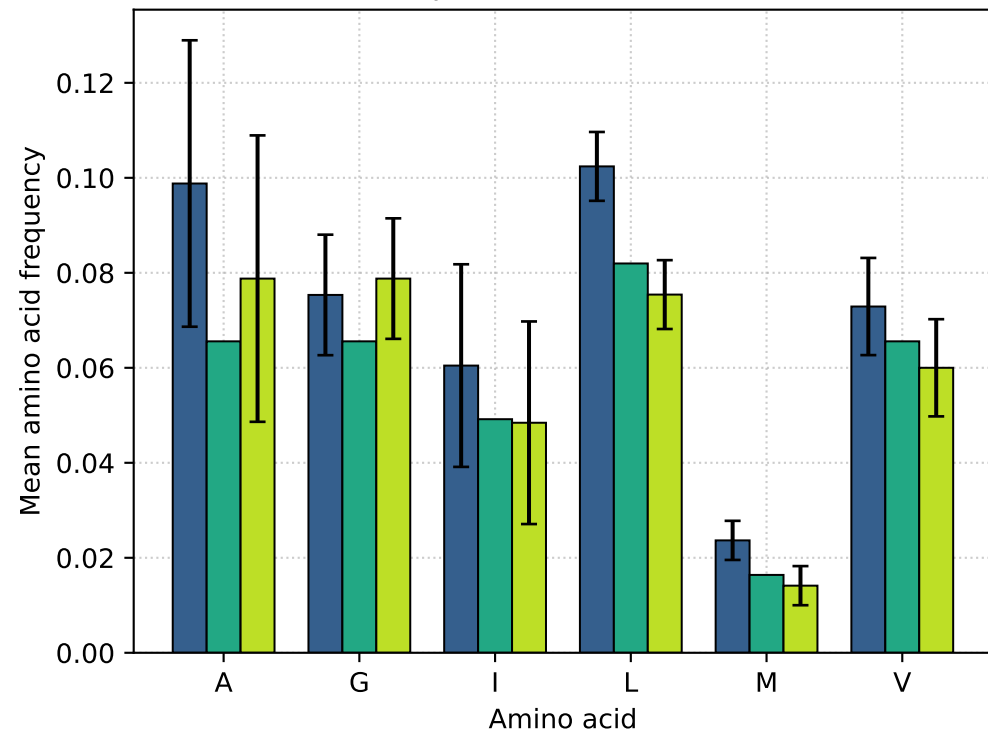


Bacteria - Mean proteome amino acid frequency for genetic code: Alternative yeast

Aliphatic amino acids



Observed
Codon
Codon+GC

Codon correlation:

- Mean Pearson:
- Coefficient: 0.55971
- p-value: 4.01511e-02

- Mean Spearman:
- Coefficient: 0.66198
- p-value: 1.70191e-02

- Mean log-MSE: 0.52546

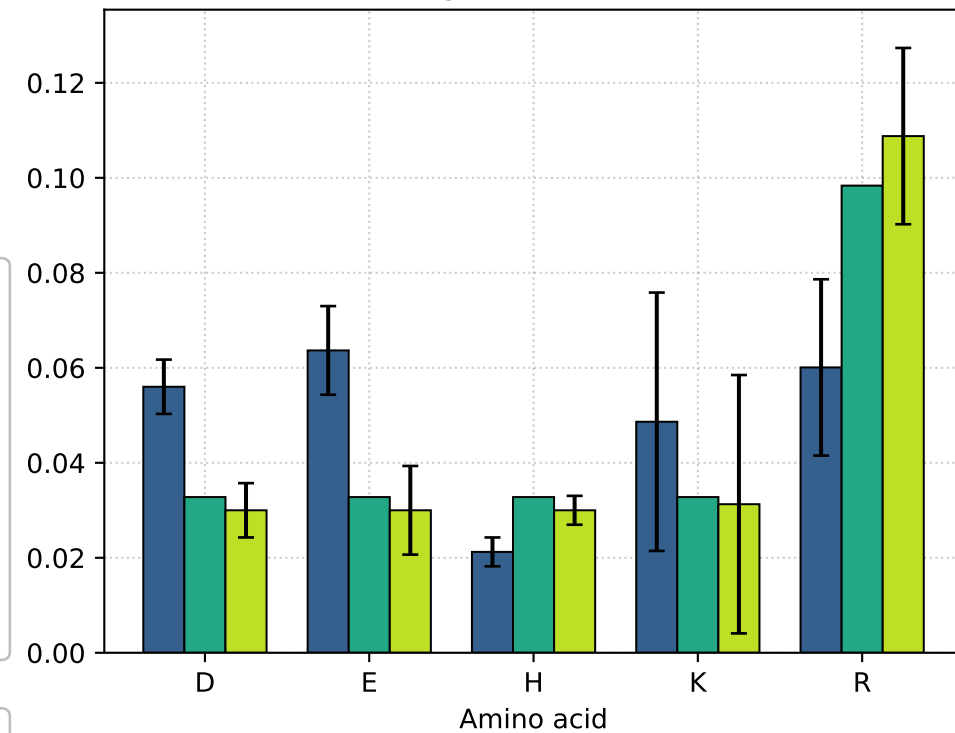
Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.65980
- p-value: 3.12215e-03

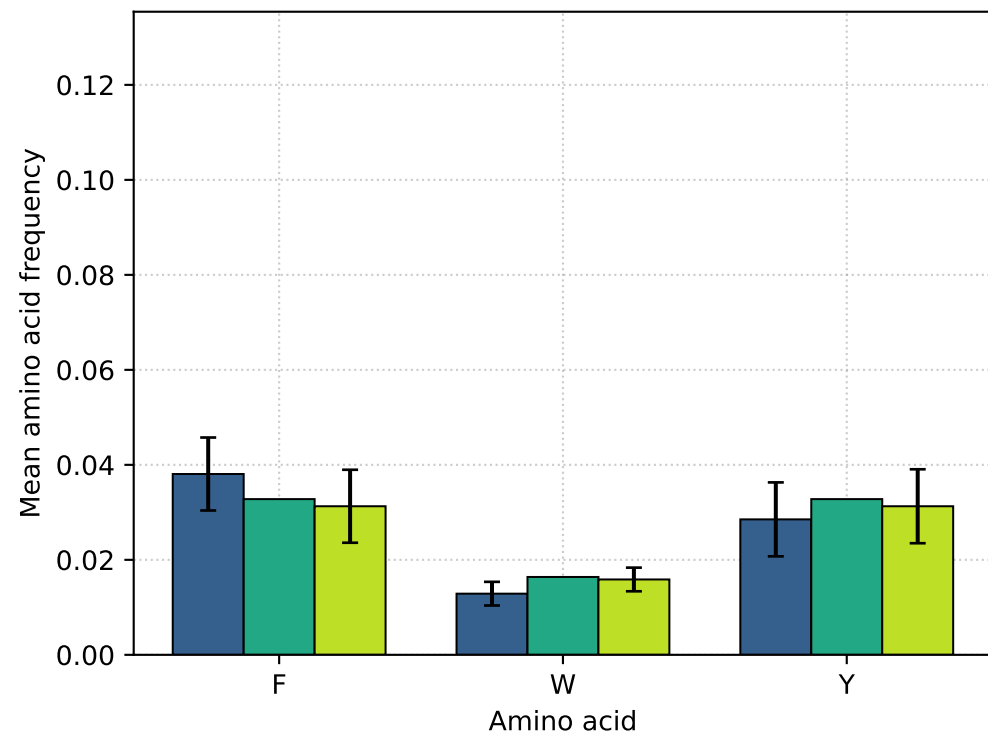
- Mean Spearman:
- Coefficient: 0.68946
- p-value: 2.37870e-03

- Mean log-MSE: 0.59265

Charged amino acids



Aromatic amino acids



Uncharged amino acids

