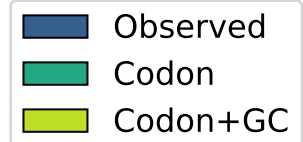
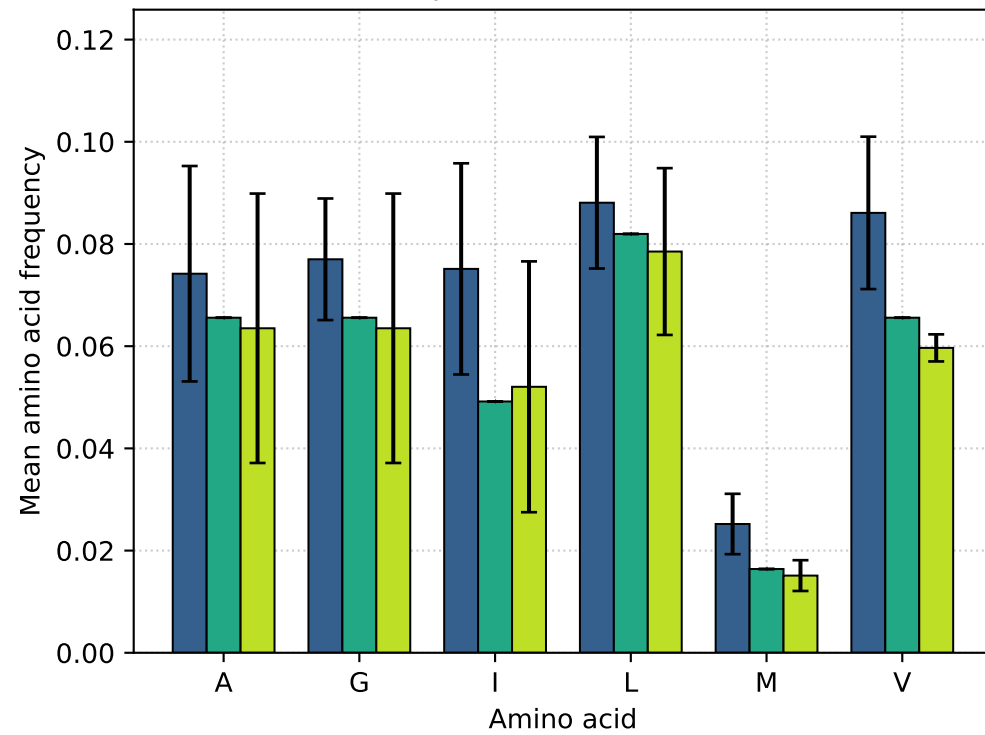


Archaea - Mean proteome amino acid frequency for genetic code: Alternative yeast

Aliphatic amino acids



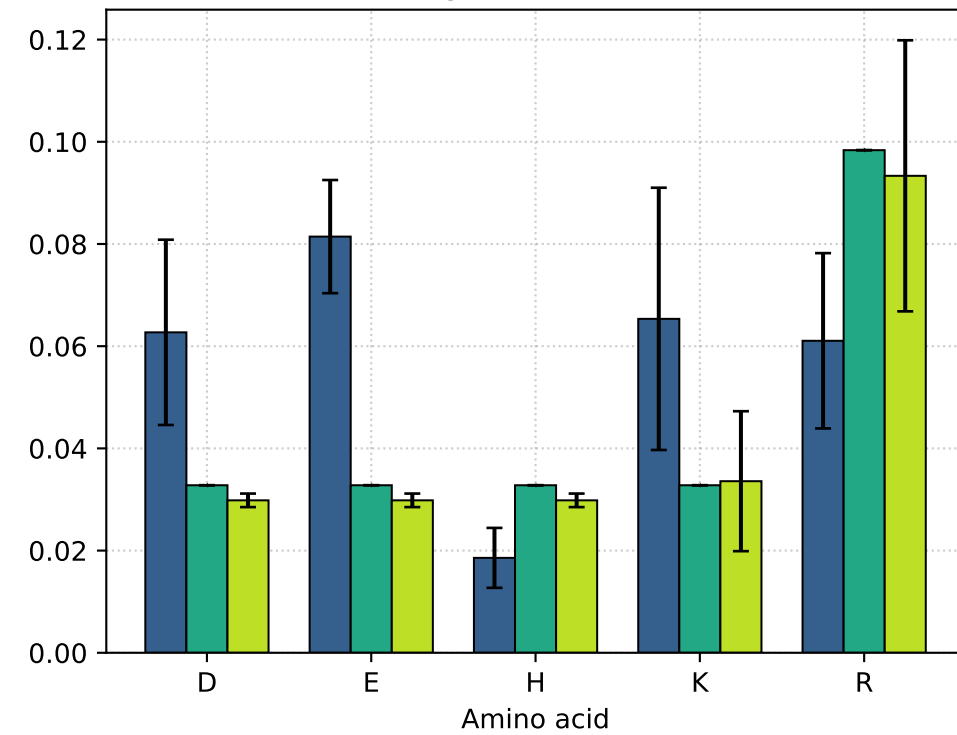
Codon correlation:
- Mean Pearson:
- Coefficient: 0.47734
- p-value: 6.66703e-02

- Mean Spearman:
- Coefficient: 0.58021
- p-value: 1.95067e-02

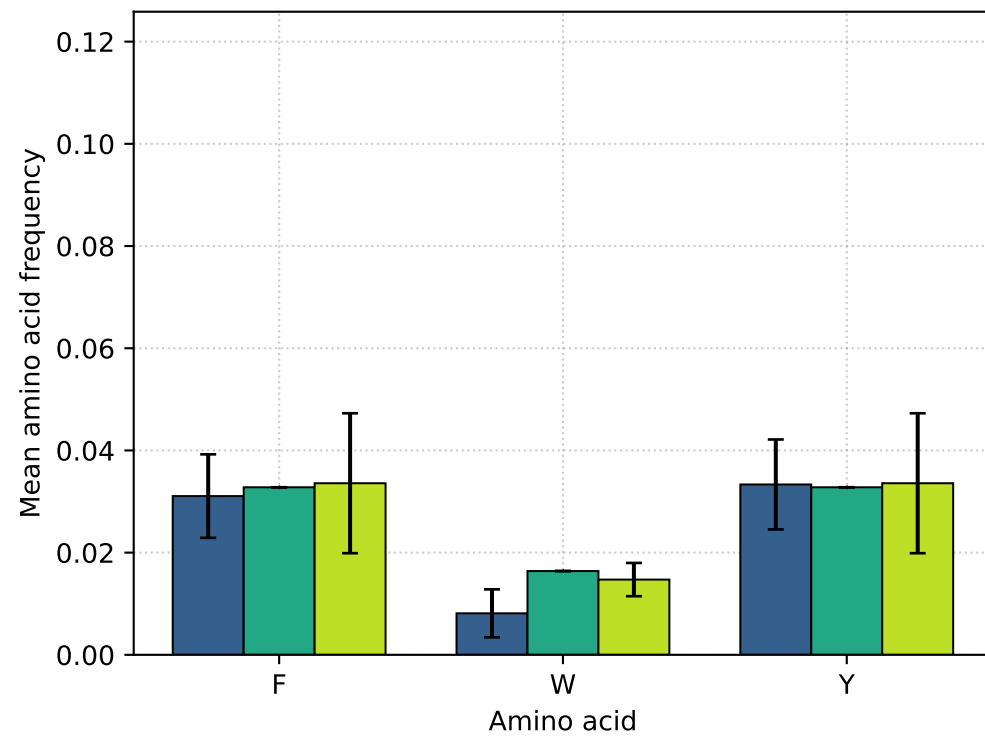
Codon + GC correlation:
- Mean Pearson:
- Coefficient: 0.53118
- p-value: 3.96324e-02

- Mean Spearman:
- Coefficient: 0.58015
- p-value: 1.78150e-02

Charged amino acids



Aromatic amino acids



Uncharged amino acids

