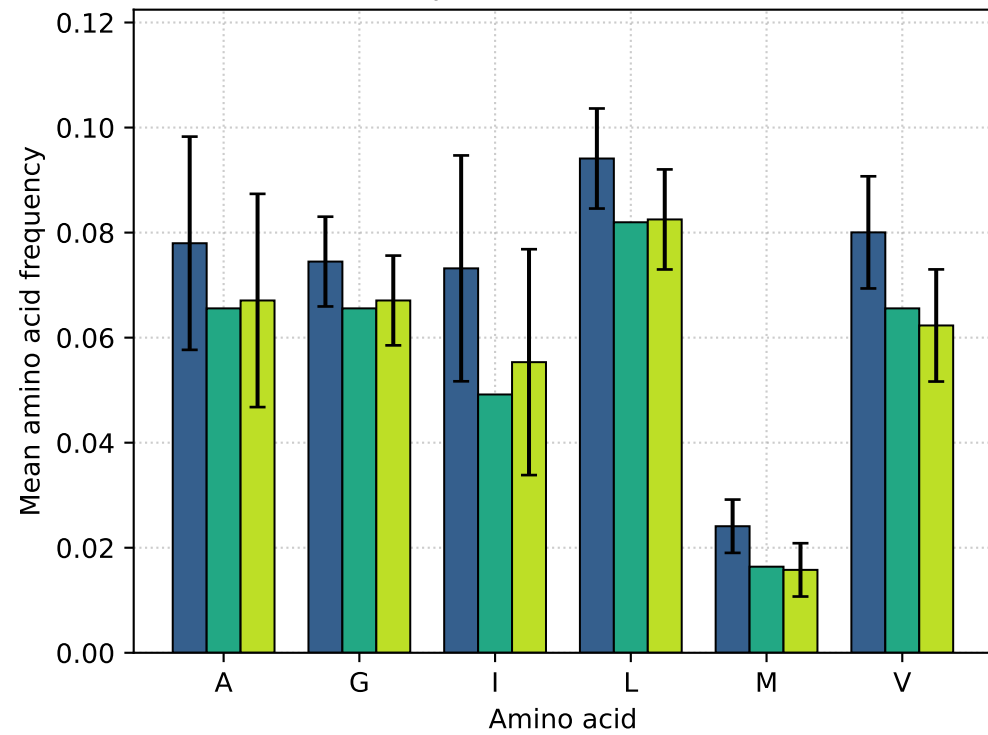


Archaea - Mean proteome amino acid frequency for genetic code: Alternative yeast

Aliphatic amino acids



Legend: Observed (dark blue), Codon (teal), Codon+GC (yellow-green)

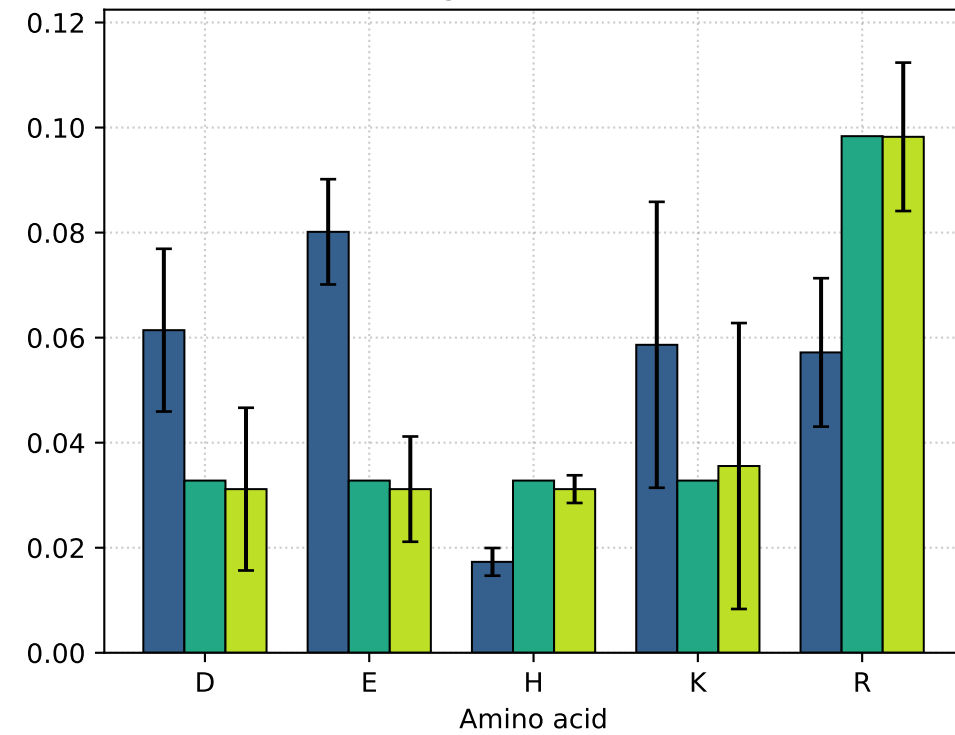
Codon correlation:

- Mean Pearson: 0.50674
- Coefficient: 0.50674
- p-value: 5.61321e-02

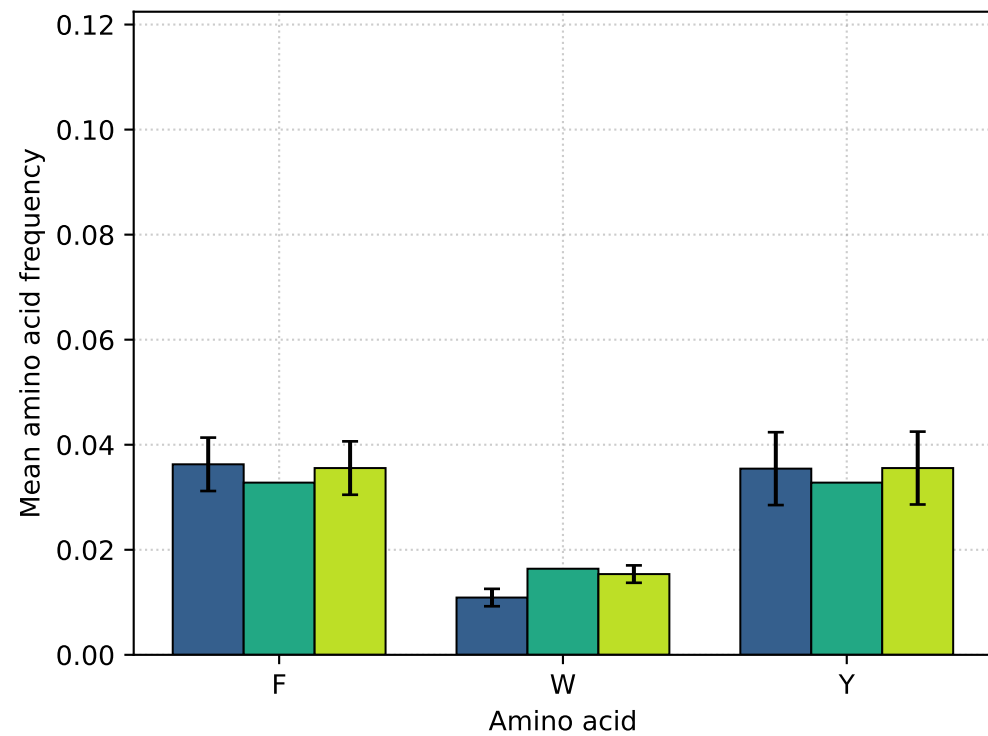
Codon + GC correlation:

- Mean Pearson: 0.59500
- Coefficient: 0.59500
- p-value: 9.25052e-03

Charged amino acids



Aromatic amino acids



Uncharged amino acids

