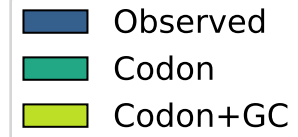
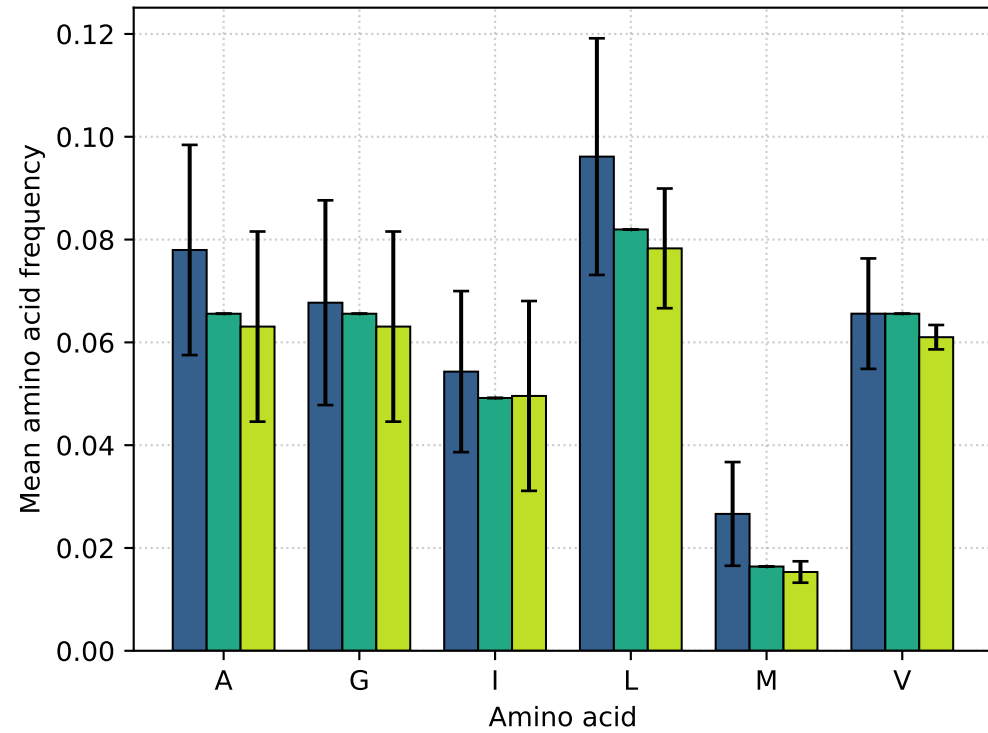


Eukaryota - Mean proteome amino acid frequency for genetic code: Alternative yeast

Aliphatic amino acids



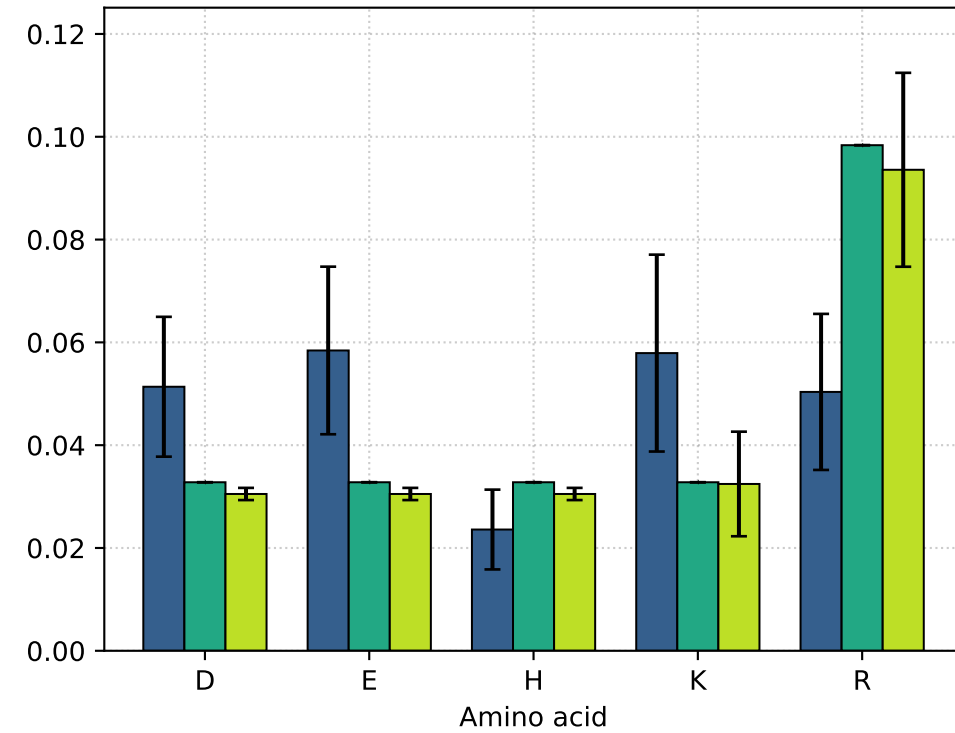
Codon correlation:
- Mean Pearson:
- Coefficient: 0.60290
- p-value: 3.18859e-02

- Mean Spearman:
- Coefficient: 0.66495
- p-value: 1.64435e-02

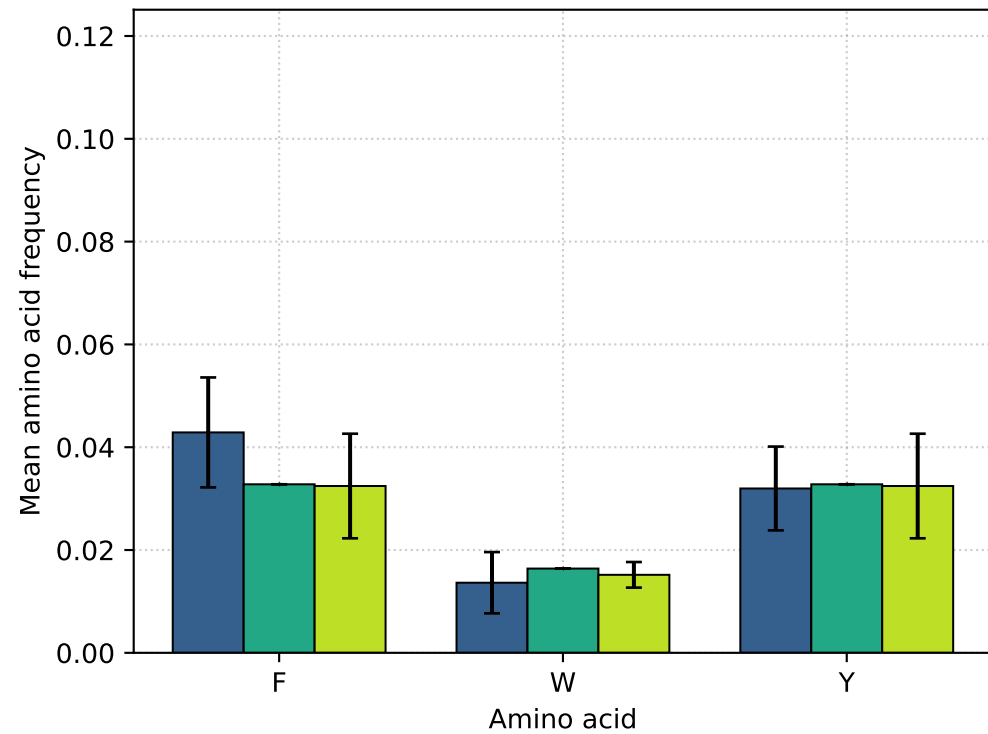
Codon + GC correlation:
- Mean Pearson:
- Coefficient: 0.63477
- p-value: 1.63994e-02

- Mean Spearman:
- Coefficient: 0.66350
- p-value: 1.07360e-02

Charged amino acids



Aromatic amino acids



Uncharged amino acids

