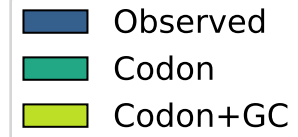
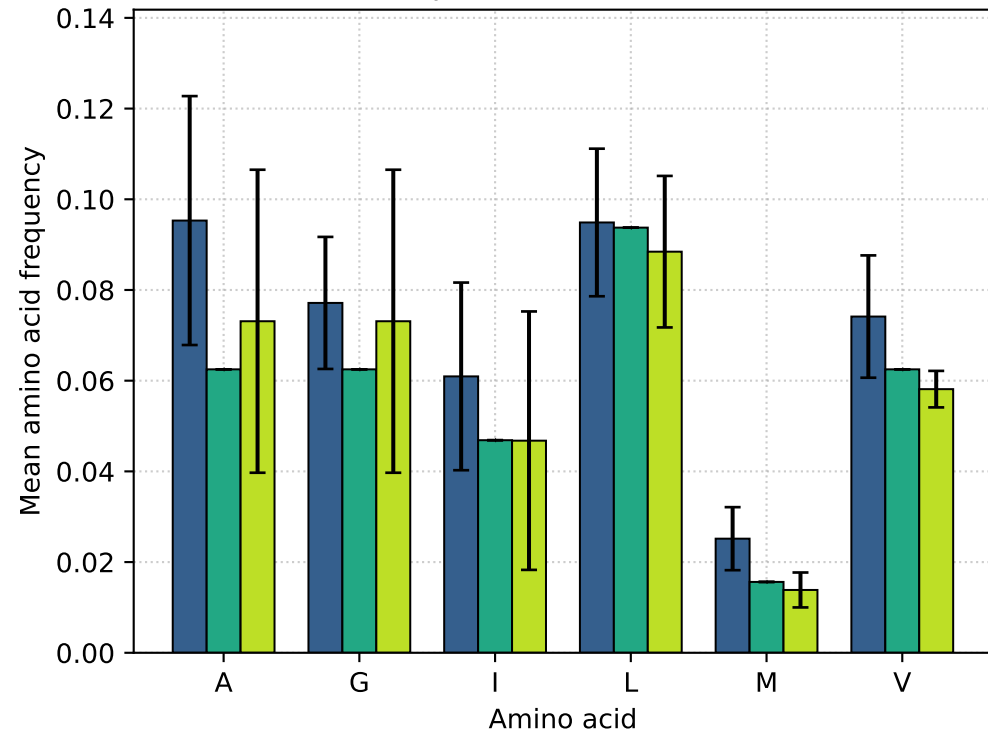


Bacteria - Mean proteome amino acid frequency for genetic code: Karyorelict nuclear

Aliphatic amino acids



Codon correlation:

- Mean Pearson: 0.51101
- Coefficient: 0.51101
- p-value: 7.25587e-02

Codon + GC correlation:

- Mean Pearson: 0.54333
- Coefficient: 0.54333
- p-value: 5.91741e-02

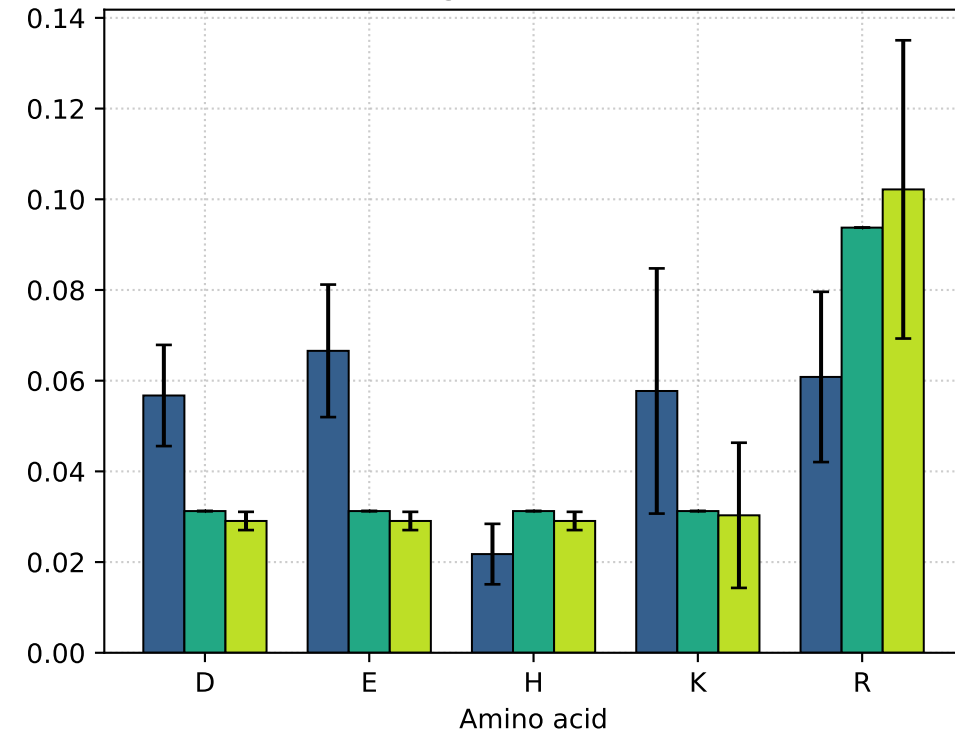
Codon + GC correlation:

- Mean Pearson: 0.58784
- Coefficient: 0.58784
- p-value: 2.73196e-02

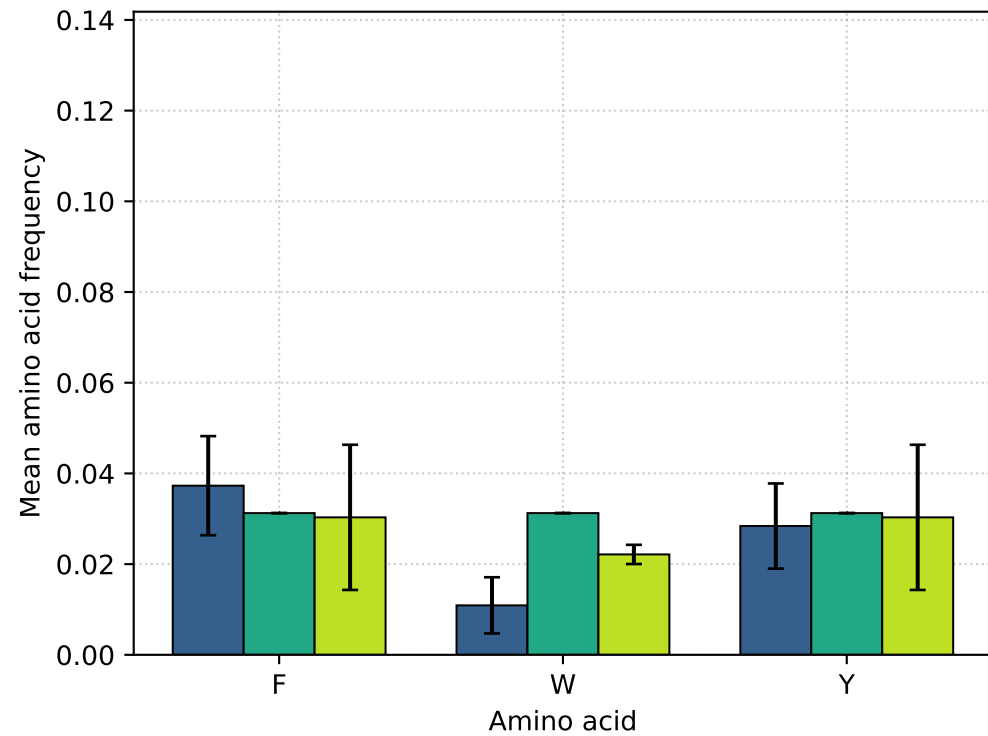
Codon correlation:

- Mean Spearman: 0.58800
- Coefficient: 0.58800
- p-value: 2.79856e-02

Charged amino acids



Aromatic amino acids



Uncharged amino acids

