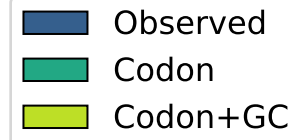
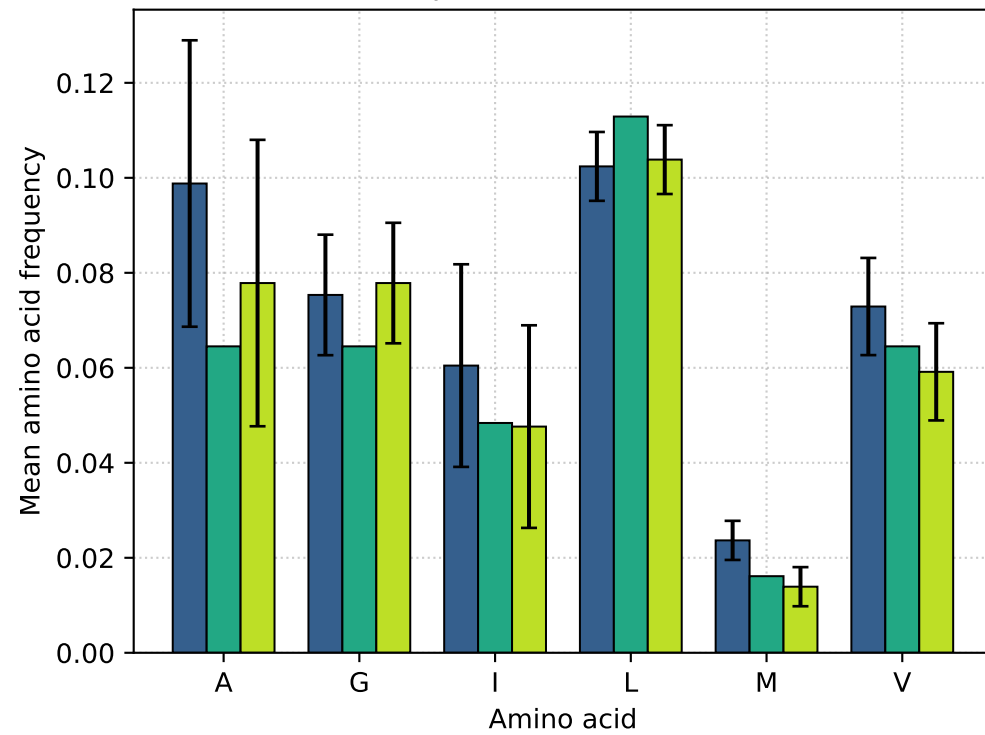


Bacteria - Mean proteome amino acid frequency for genetic code: Chlorophycean mitochondrial

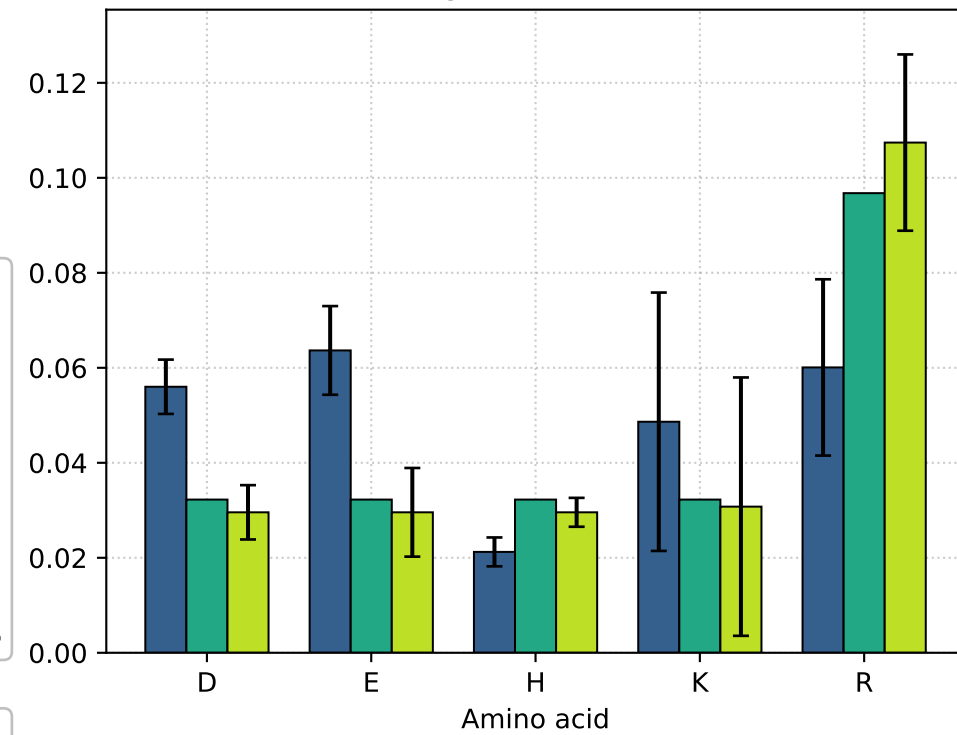
Aliphatic amino acids



Codon correlation:

- Mean Pearson:
- Coefficient: 0.64120
- p-value: 2.02593e-02
- Mean Spearman:
- Coefficient: 0.68217
- p-value: 1.36750e-02
- Mean log-MSE: 0.49845

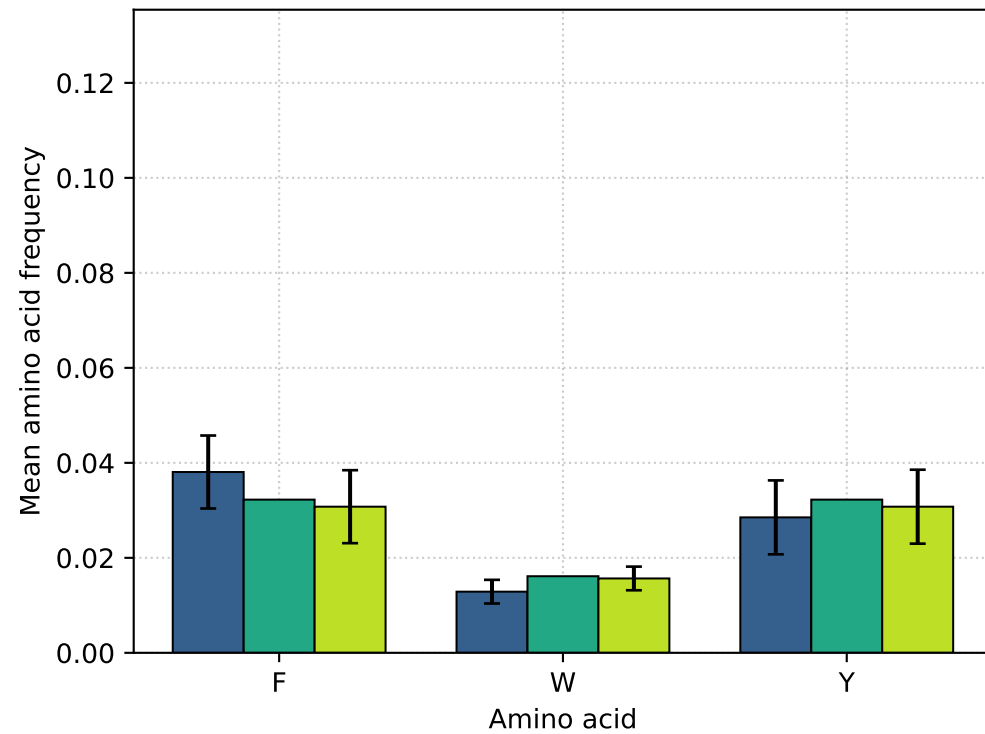
Charged amino acids



Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.72195
- p-value: 8.24307e-04
- Mean Spearman:
- Coefficient: 0.70437
- p-value: 1.93224e-03
- Mean log-MSE: 0.55732

Aromatic amino acids



Uncharged amino acids

