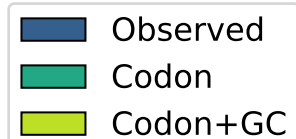
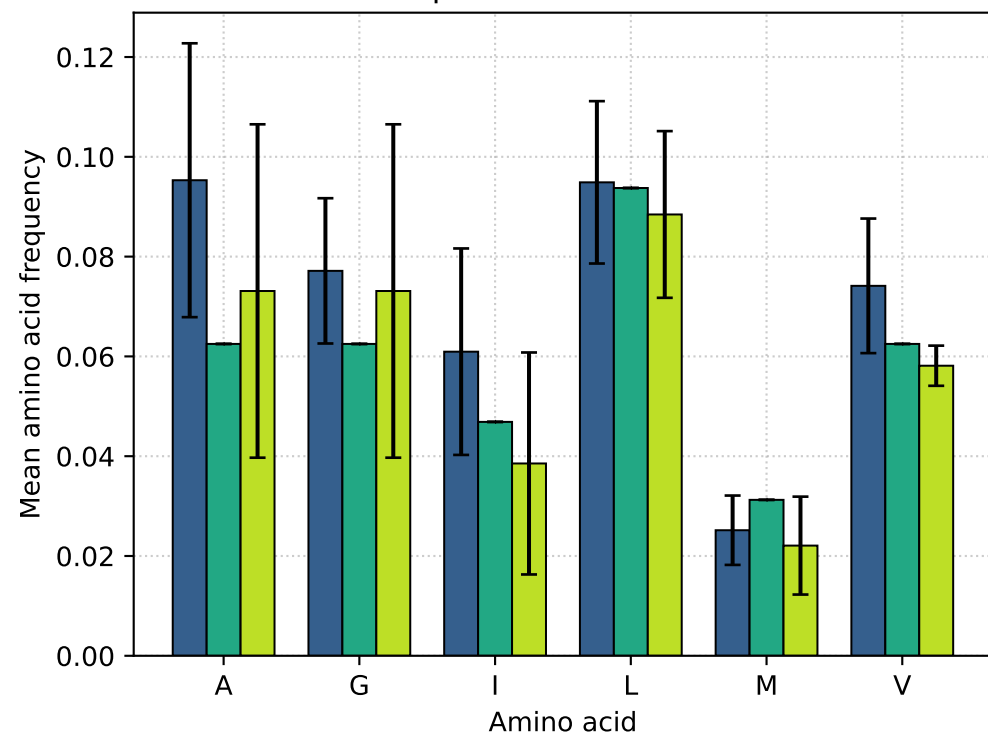


Bacteria - Mean proteome amino acid frequency for genetic code: Rhabdopleuridae mitochondrial

Aliphatic amino acids



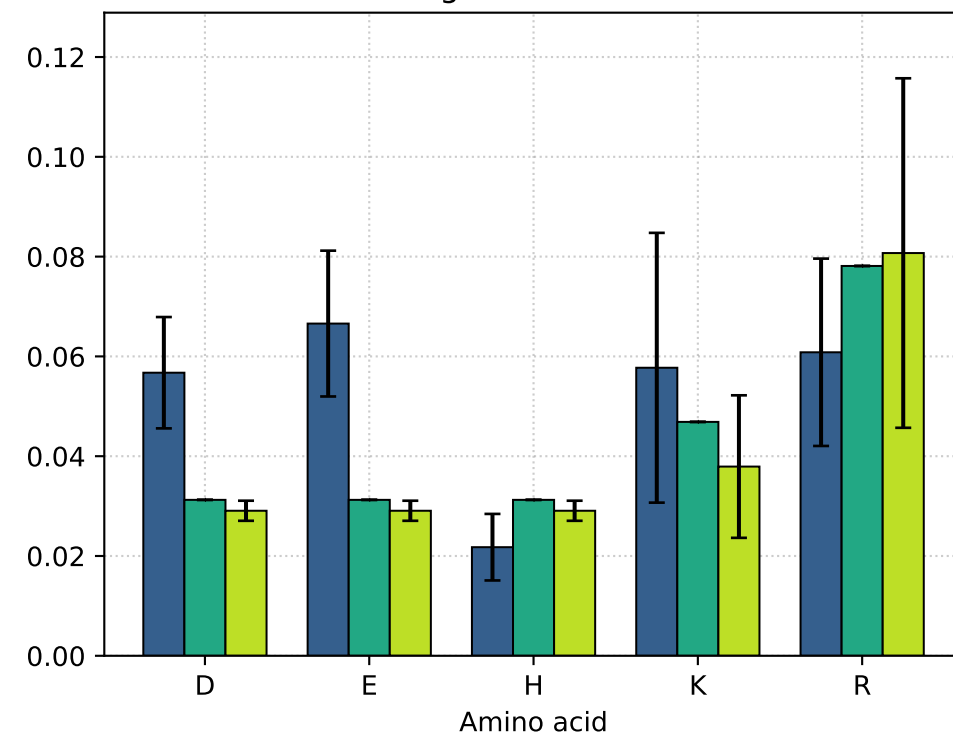
Codon correlation:
- Mean Pearson:
- Coefficient: 0.53241
- p-value: 4.26288e-02

- Mean Spearman:
- Coefficient: 0.60752
- p-value: 2.52550e-02

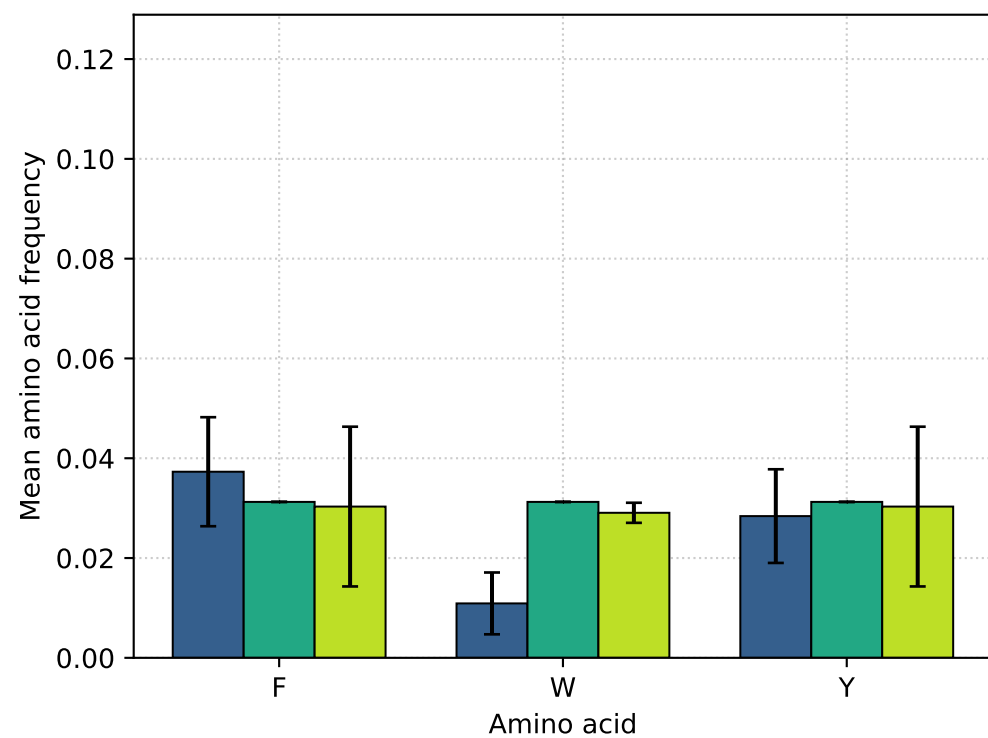
Codon + GC correlation:
- Mean Pearson:
- Coefficient: 0.59443
- p-value: 2.34364e-02

- Mean Spearman:
- Coefficient: 0.59868
- p-value: 2.11566e-02

Charged amino acids



Aromatic amino acids



Uncharged amino acids

