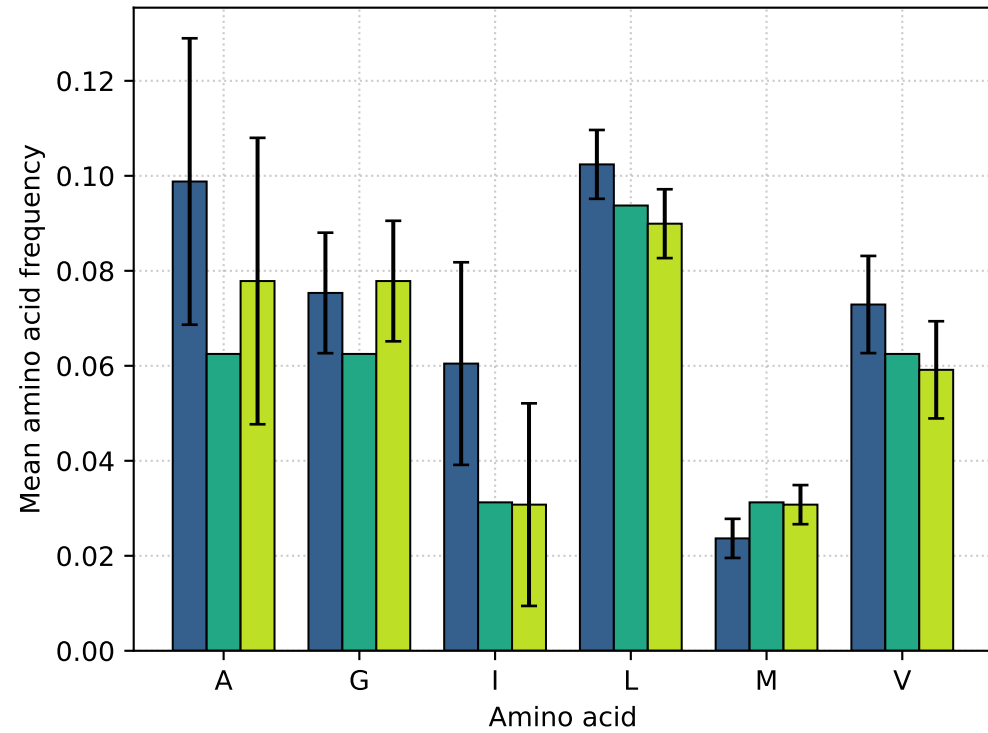


Bacteria - Mean proteome amino acid frequency for genetic code: Invertebrate mitochondrial

Aliphatic amino acids



Observed
Codon
Codon+GC

Codon correlation:

- Mean Pearson:
- Coefficient: 0.52255
- p-value: 4.36548e-02

- Mean Spearman:
- Coefficient: 0.59495
- p-value: 3.35172e-02

- Mean log-MSE: 0.61808

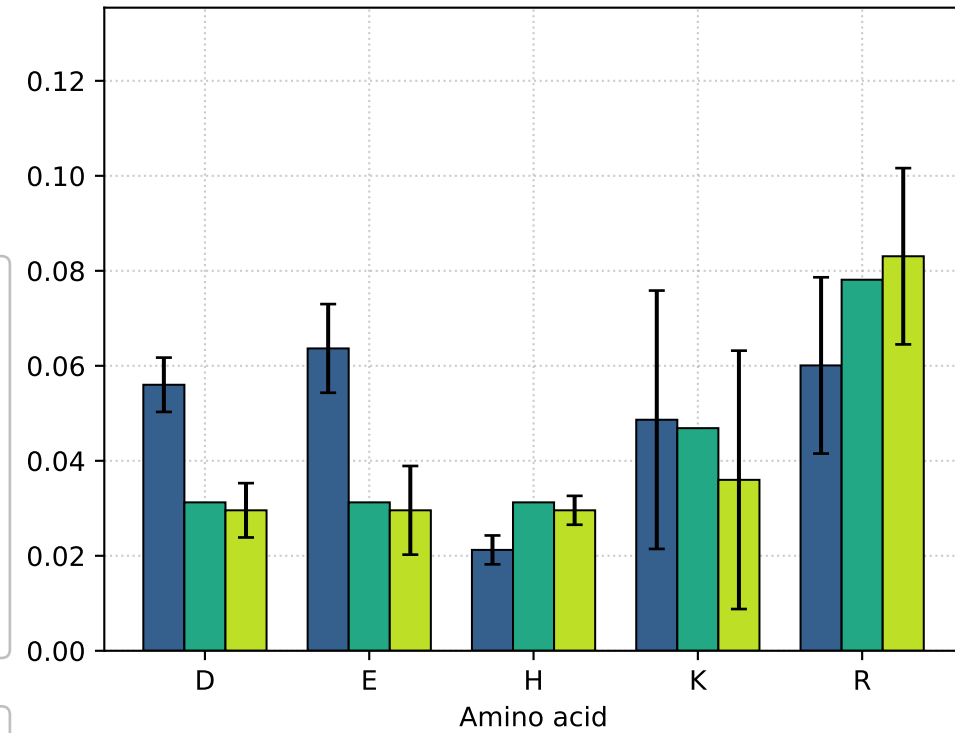
Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.64371
- p-value: 7.14043e-03

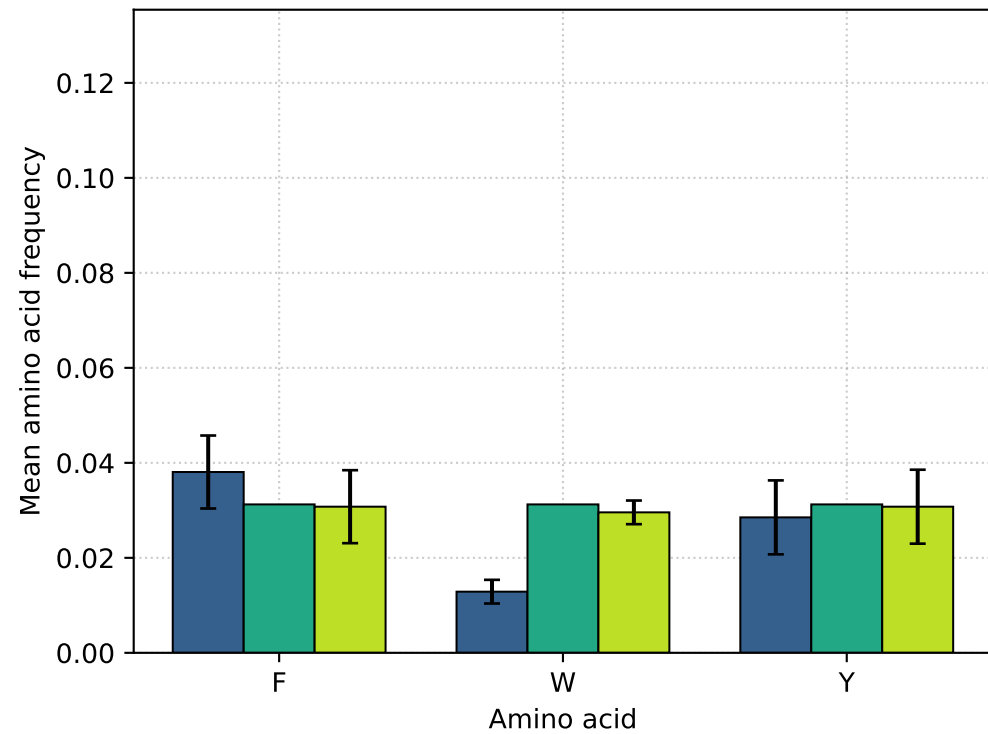
- Mean Spearman:
- Coefficient: 0.61238
- p-value: 1.18085e-02

- Mean log-MSE: 0.64686

Charged amino acids



Aromatic amino acids



Uncharged amino acids

