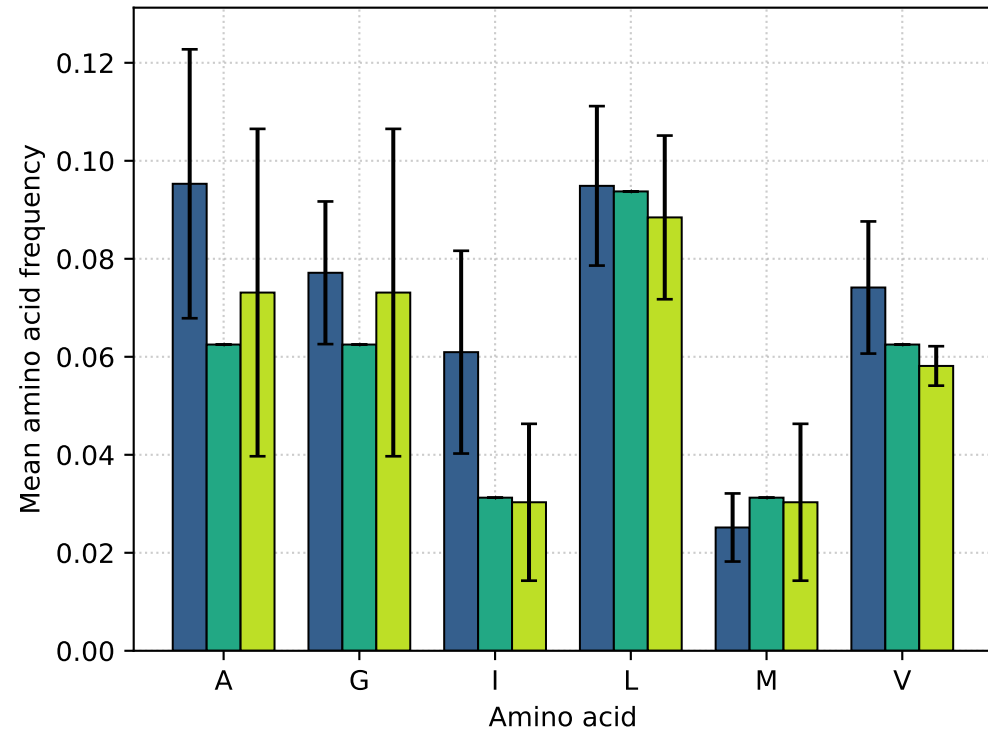


Bacteria - Mean proteome amino acid frequency for genetic code: Invertebrate mitochondrial

Aliphatic amino acids



Observed
Codon
Codon+GC

Codon correlation:

- Mean Pearson:
- Coefficient: 0.47132
- p-value: 7.12222e-02

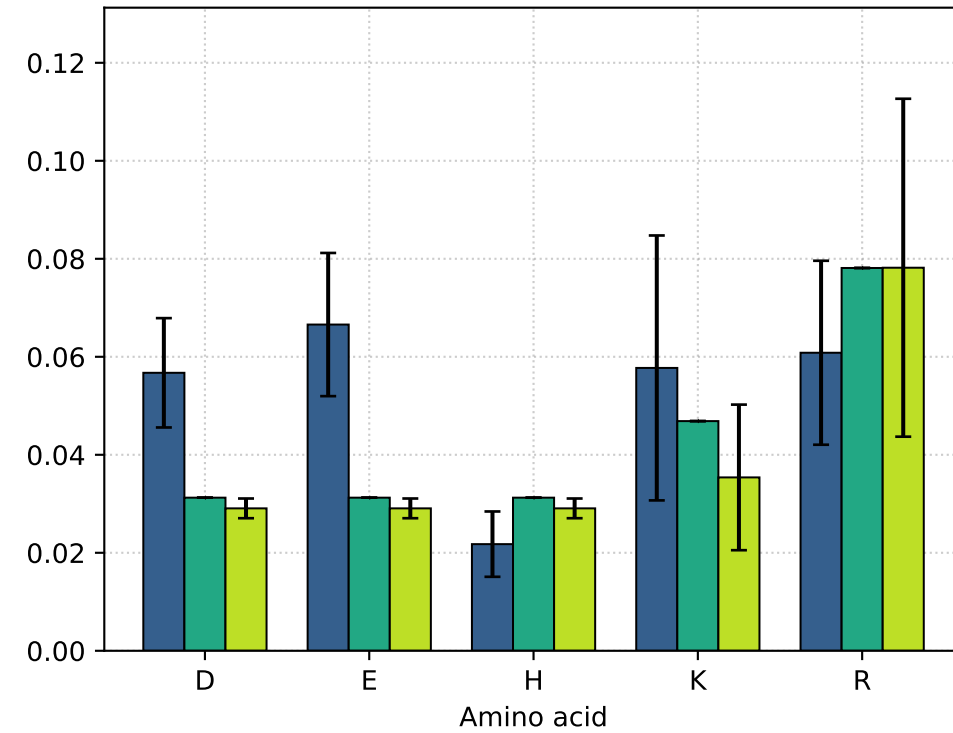
- Mean Spearman:
- Coefficient: 0.56582
- p-value: 4.01053e-02

Codon + GC correlation:

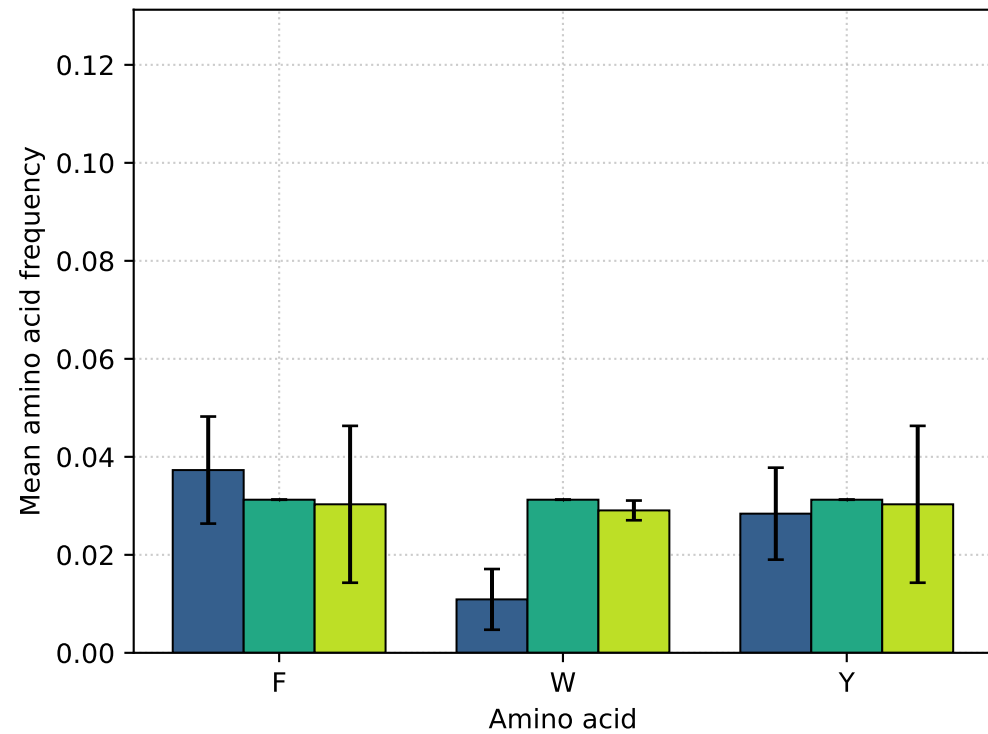
- Mean Pearson:
- Coefficient: 0.55832
- p-value: 3.75033e-02

- Mean Spearman:
- Coefficient: 0.55104
- p-value: 3.69038e-02

Charged amino acids



Aromatic amino acids



Uncharged amino acids

