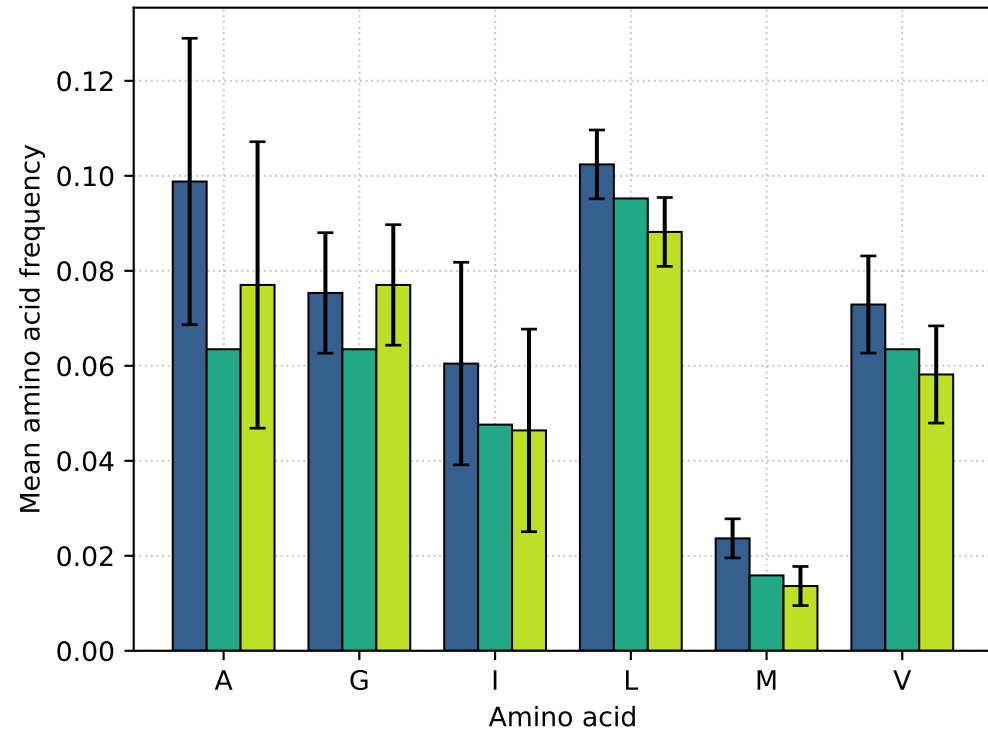


Bacteria - Mean proteome amino acid frequency for genetic code: Alternative flatworm mitochondrial

Aliphatic amino acids



Observed
Codon
Codon+GC

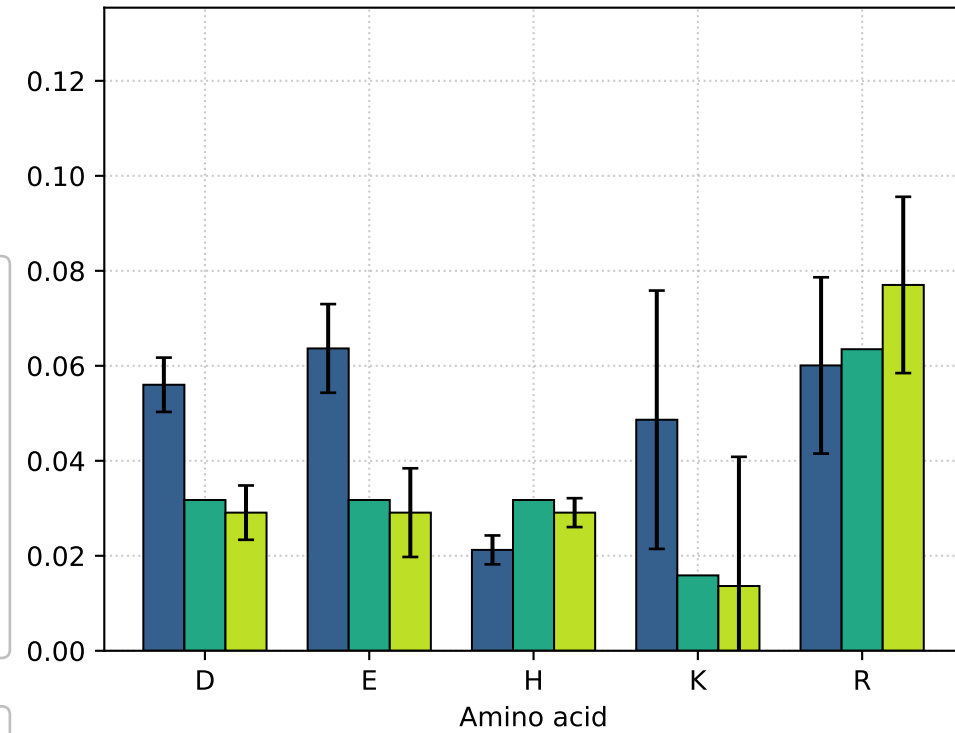
Codon correlation:

- Mean Pearson:
- Coefficient: 0.50953
- p-value: 4.75233e-02

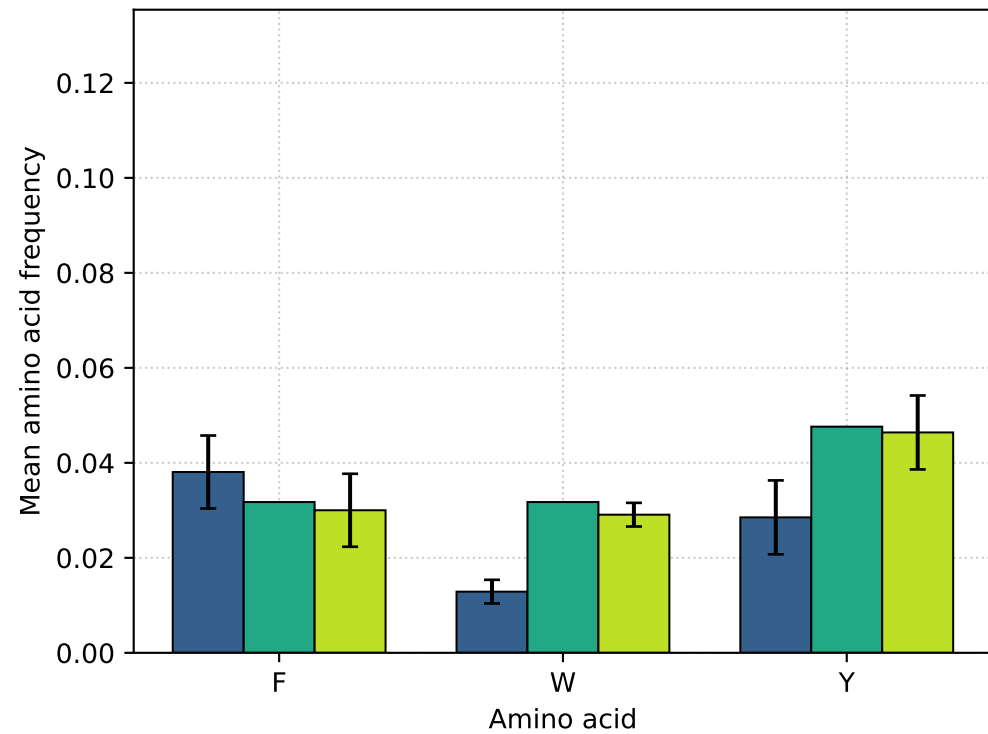
- Mean Spearman:
- Coefficient: 0.56932
- p-value: 4.31987e-02

- Mean log-MSE: 0.72574

Charged amino acids



Aromatic amino acids



Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.60874
- p-value: 1.56253e-02

- Mean Spearman:
- Coefficient: 0.57998
- p-value: 3.16654e-02

- Mean log-MSE: 0.78024

Uncharged amino acids

