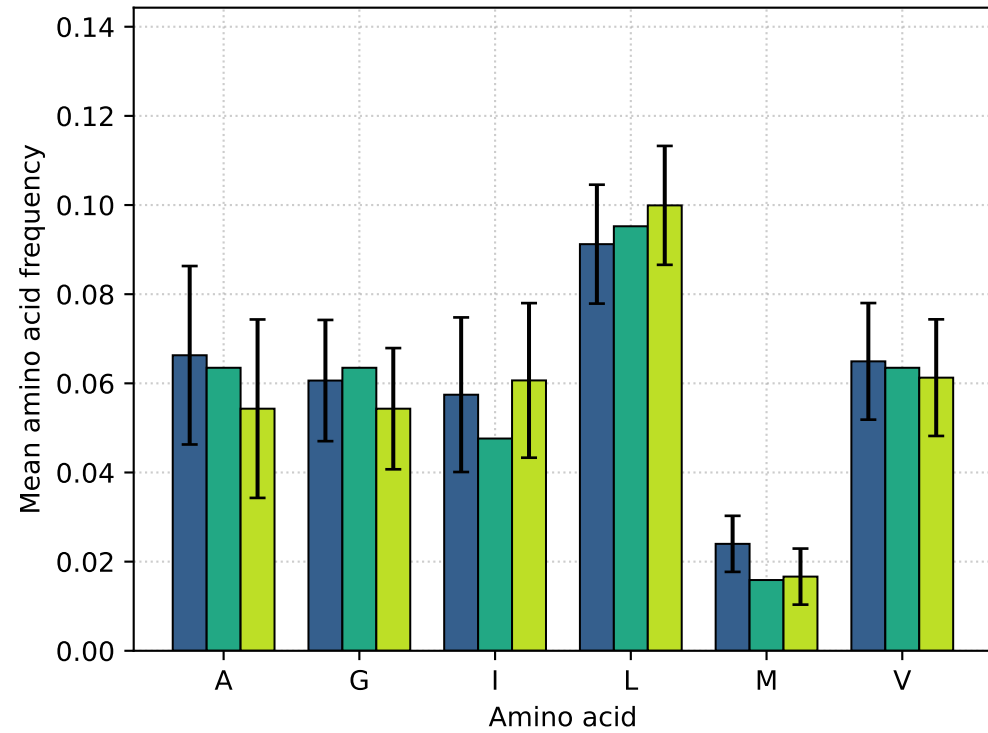


Viruses - Mean proteome amino acid frequency for genetic code: Alternative flatworm mitochondrial

Aliphatic amino acids



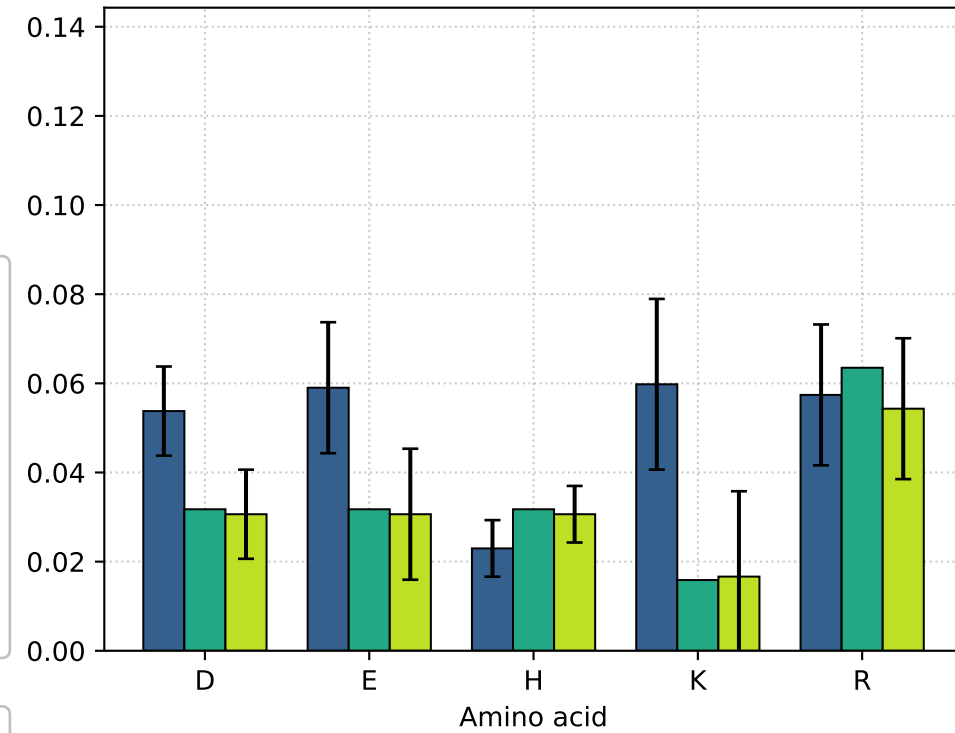
Observed
Codon
Codon+GC

Codon correlation:
- Mean Pearson:
- Coefficient: 0.57486
- p-value: 4.83336e-02

- Mean Spearman:
- Coefficient: 0.57039
- p-value: 5.20517e-02

- Mean log-MSE: 0.57843

Charged amino acids

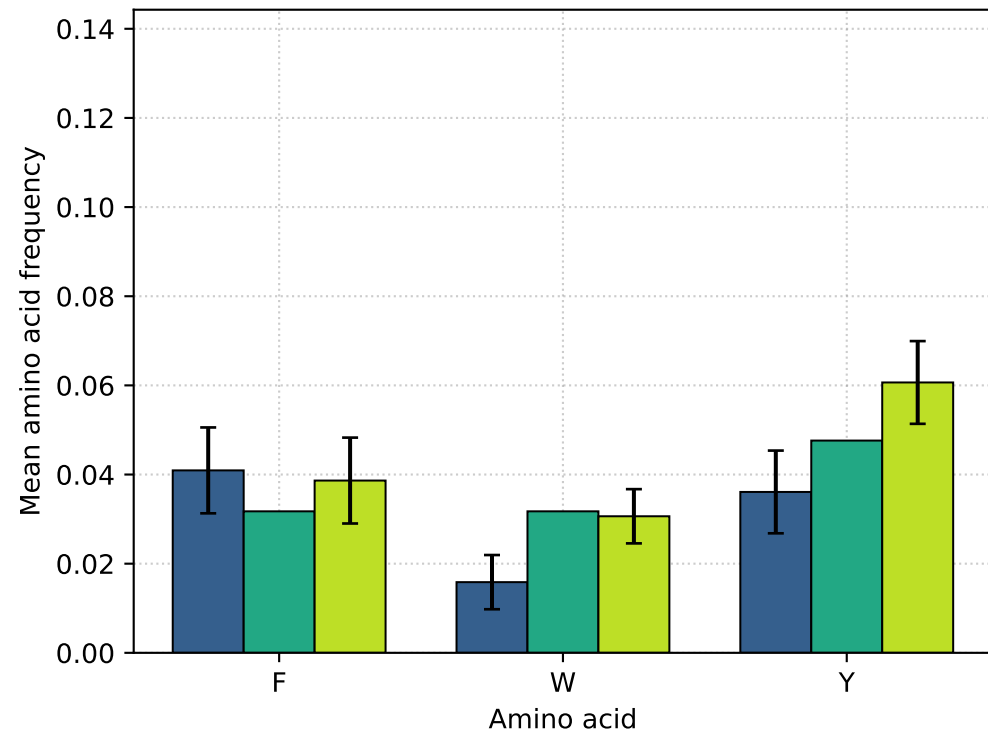


Codon + GC correlation:
- Mean Pearson:
- Coefficient: 0.60826
- p-value: 2.67356e-02

- Mean Spearman:
- Coefficient: 0.56398
- p-value: 4.12055e-02

- Mean log-MSE: 0.56747

Aromatic amino acids



Uncharged amino acids

