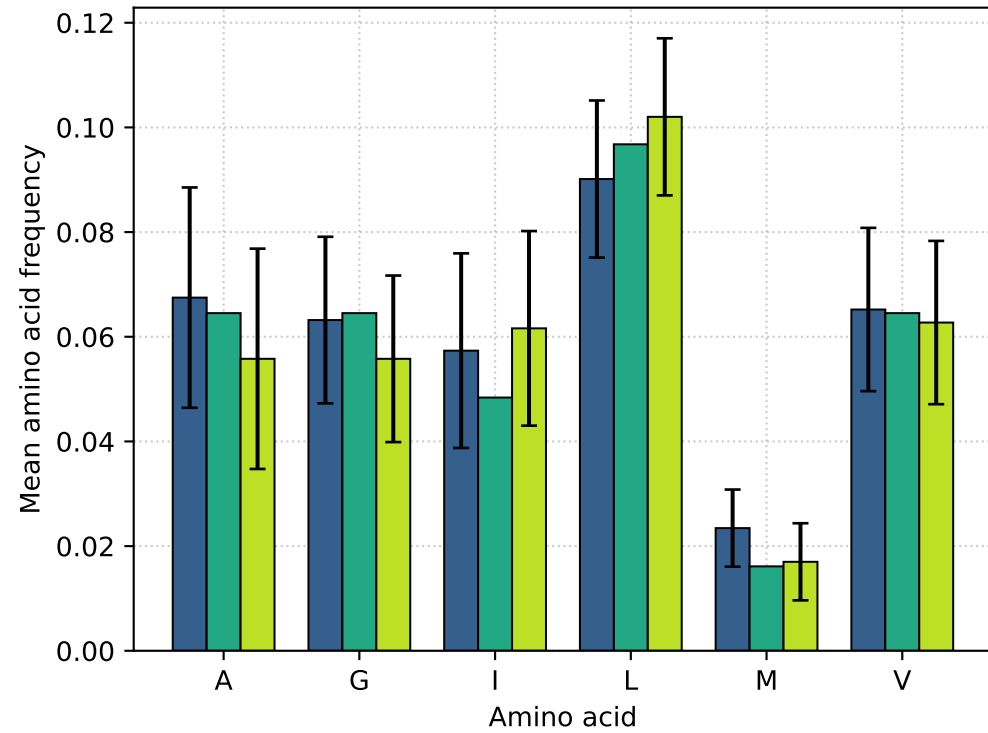


Viruses - Mean proteome amino acid frequency for genetic code: Euplotid nuclear

Aliphatic amino acids



Observed
Codon
Codon+GC

Codon correlation:

- Mean Pearson:
- Coefficient: 0.60204
- p-value: 4.51782e-02

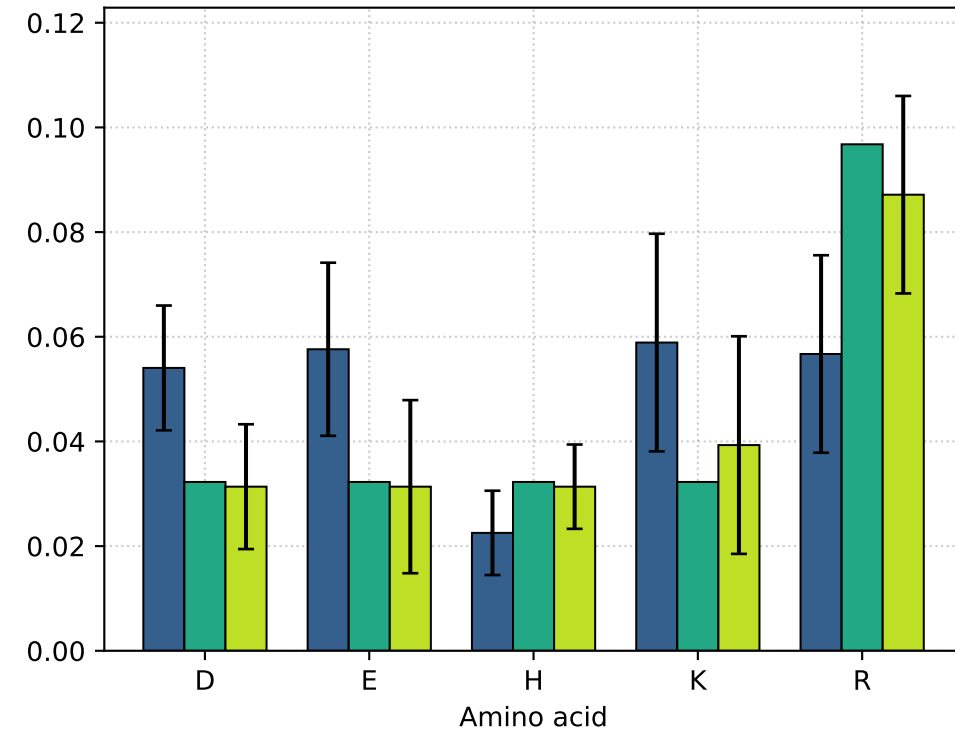
- Mean Spearman:
- Coefficient: 0.59930
- p-value: 4.31056e-02

Codon + GC correlation:

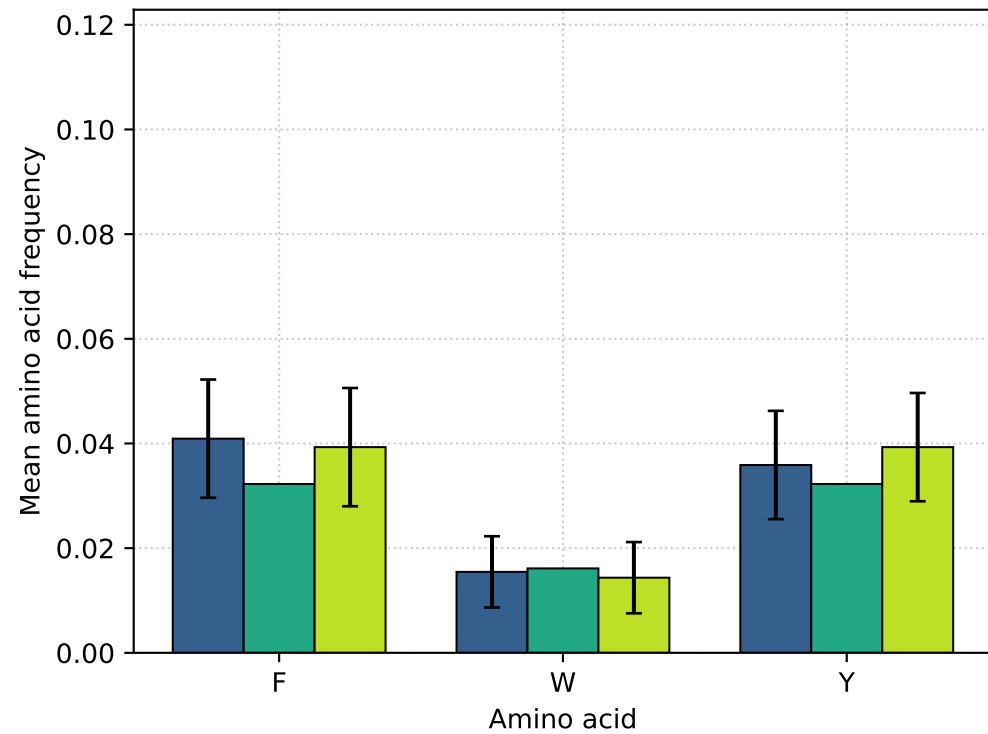
- Mean Pearson:
- Coefficient: 0.67101
- p-value: 1.27932e-02

- Mean Spearman:
- Coefficient: 0.62561
- p-value: 2.01532e-02

Charged amino acids



Aromatic amino acids



Uncharged amino acids

