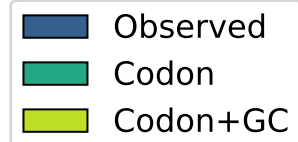
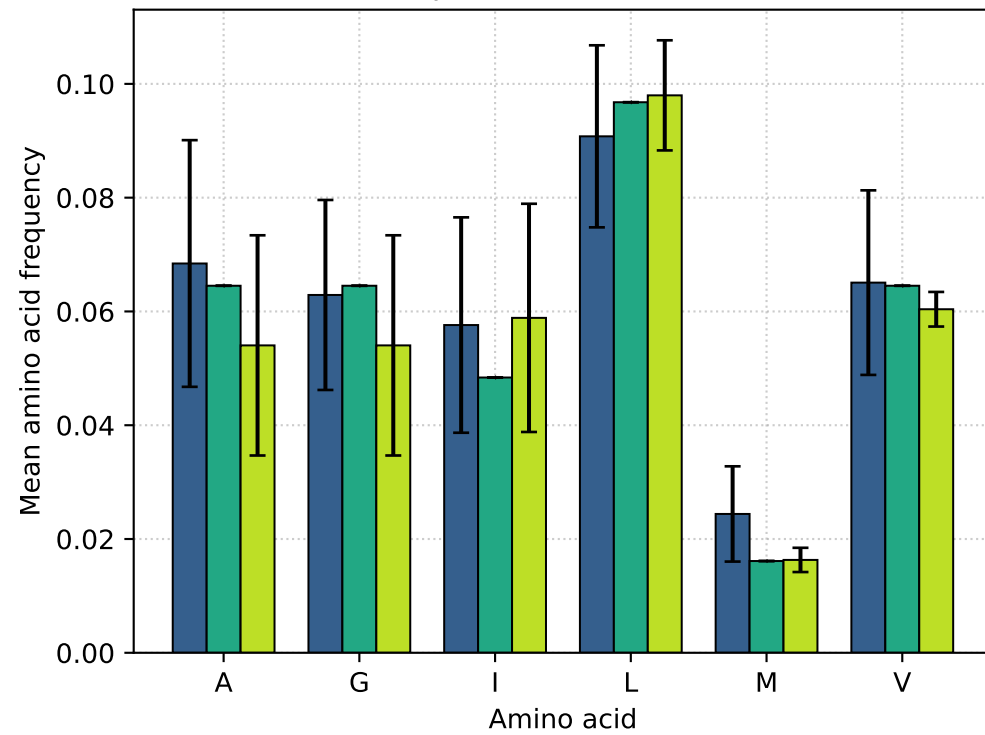


Viruses - Mean proteome amino acid frequency for genetic code: Euplotid nuclear

Aliphatic amino acids



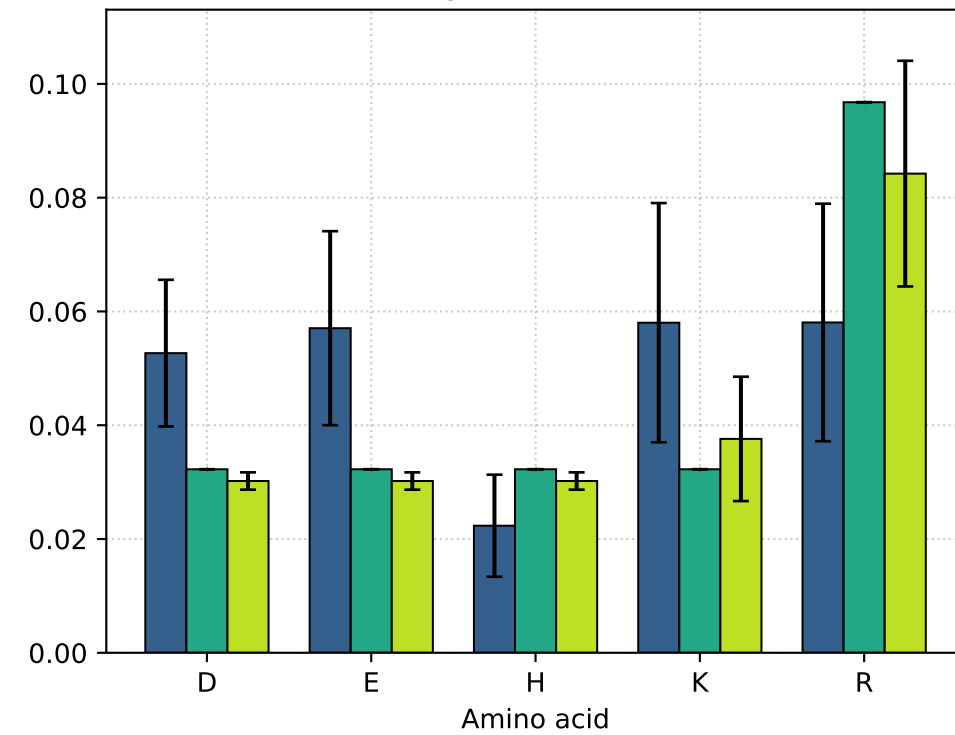
Codon correlation:
- Mean Pearson:
- Coefficient: 0.61009
- p-value: 4.50142e-02

- Mean Spearman:
- Coefficient: 0.60591
- p-value: 4.45784e-02

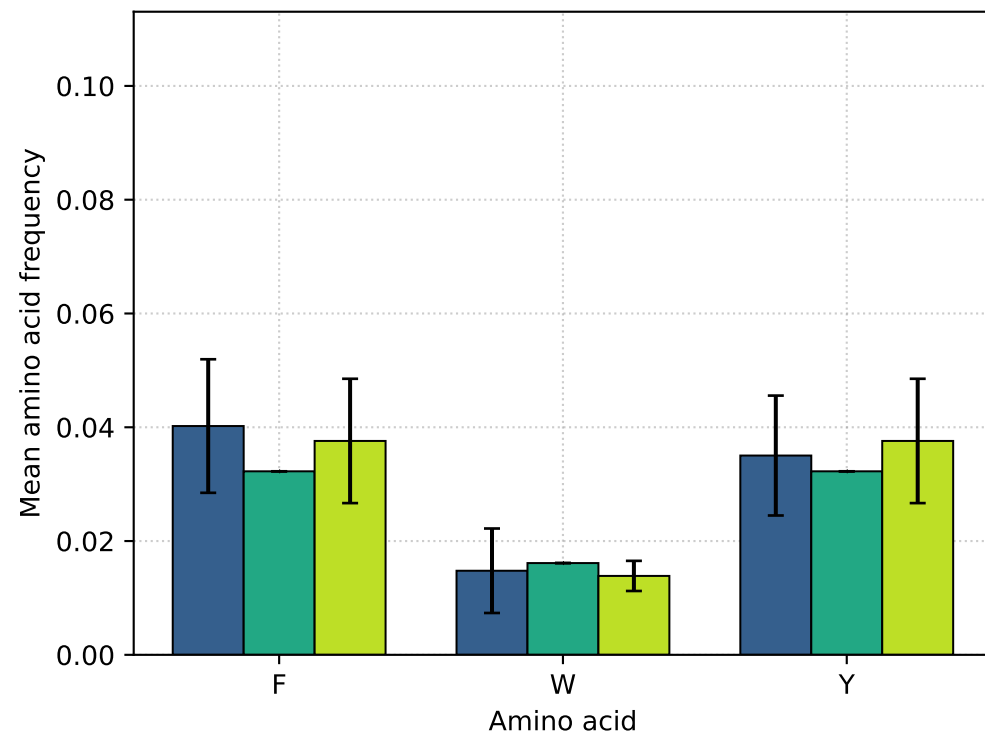
Codon + GC correlation:
- Mean Pearson:
- Coefficient: 0.68057
- p-value: 1.26146e-02

- Mean Spearman:
- Coefficient: 0.63496
- p-value: 1.97789e-02

Charged amino acids



Aromatic amino acids



Uncharged amino acids

