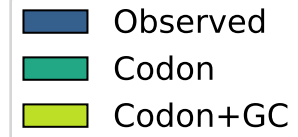
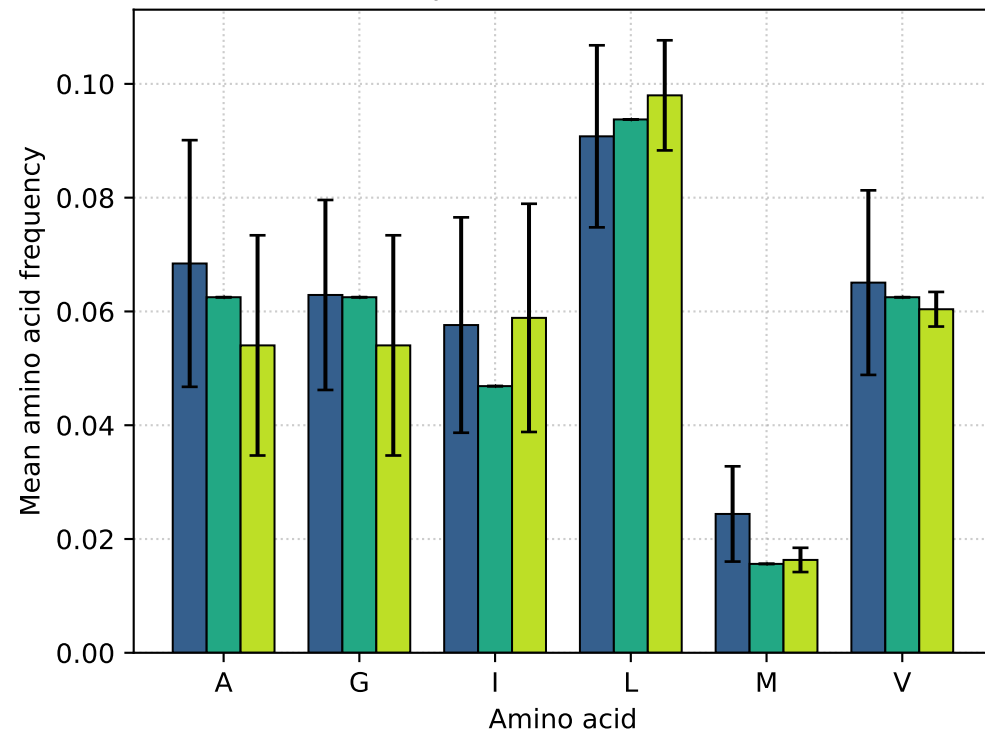


Viruses - Mean proteome amino acid frequency for genetic code: Condyllostoma nuclear

Aliphatic amino acids



Codon correlation:

- Mean Pearson:
- Coefficient: 0.58952
- p-value: 5.58668e-02

- Mean Spearman:

- Coefficient: 0.58500
- p-value: 5.81717e-02

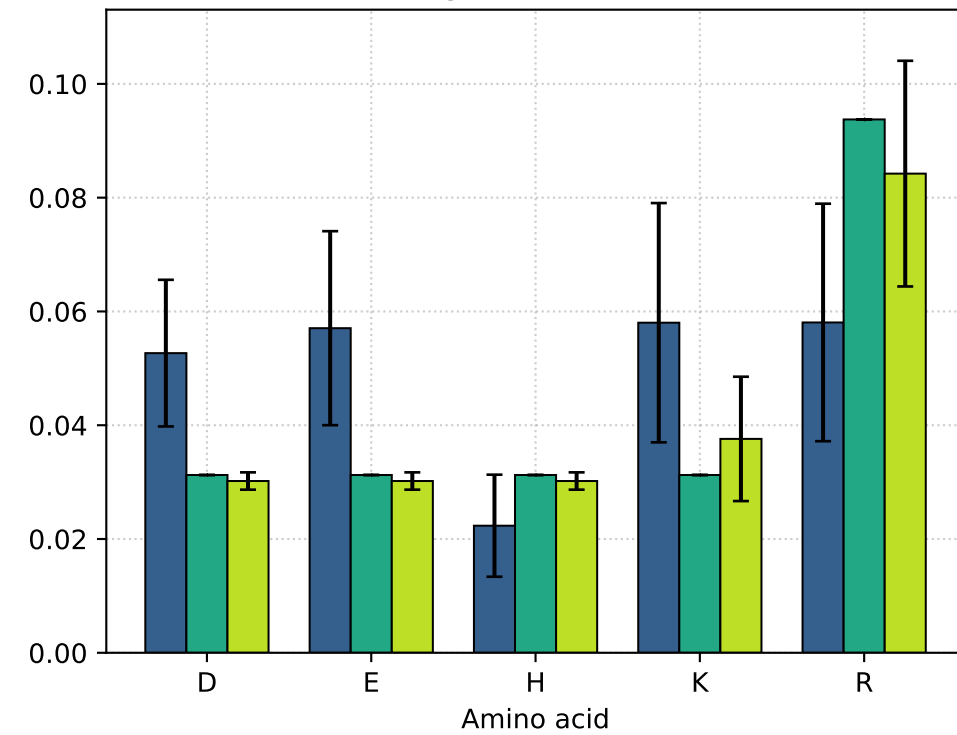
Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.69532
- p-value: 1.02925e-02

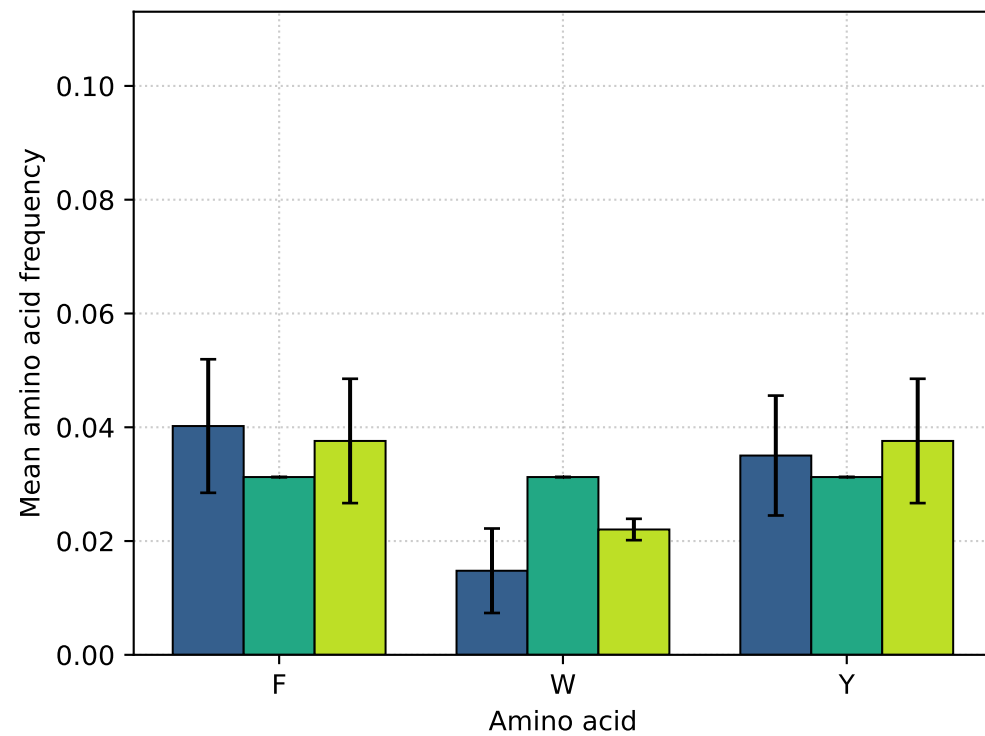
- Mean Spearman:

- Coefficient: 0.66886
- p-value: 1.23321e-02

Charged amino acids



Aromatic amino acids



Uncharged amino acids

