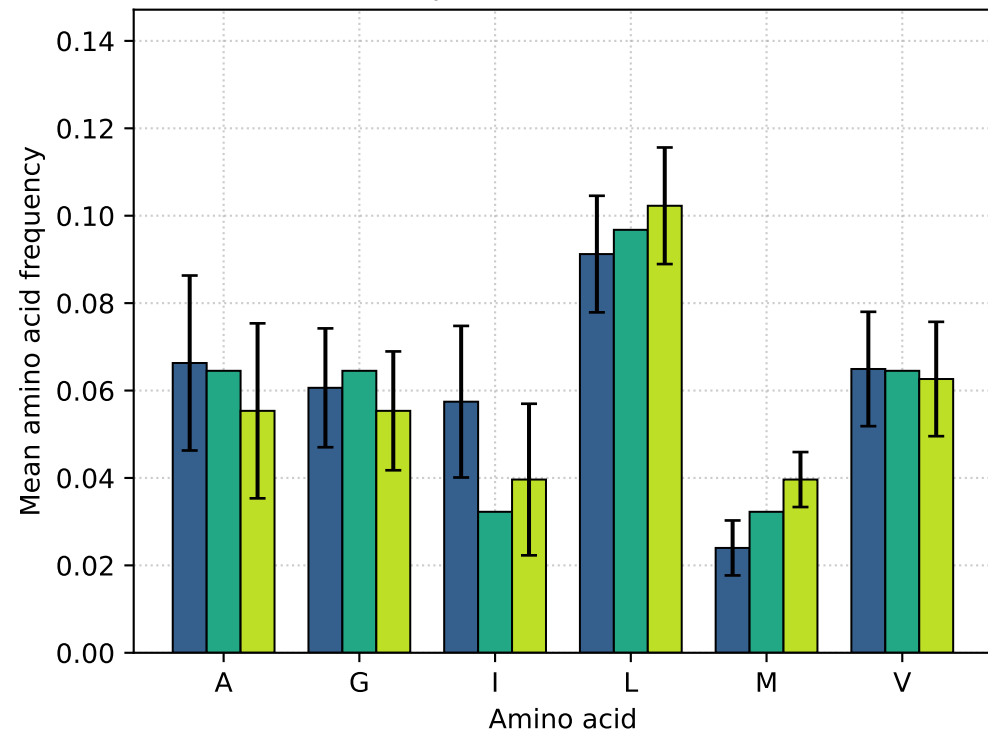
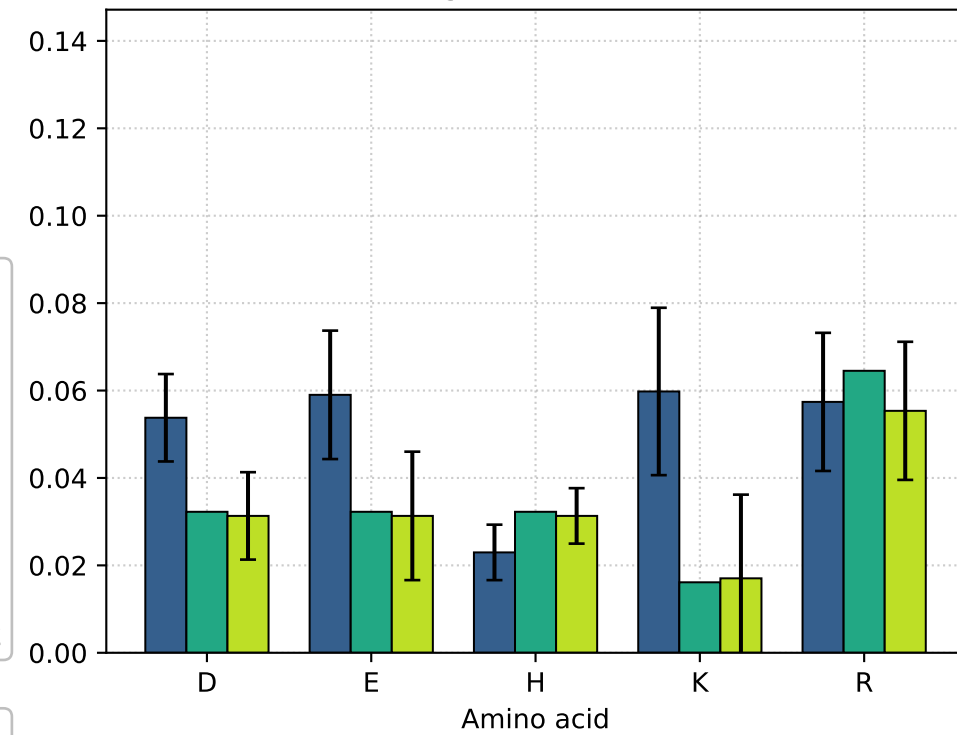


Viruses - Mean proteome amino acid frequency for genetic code: Trematode mitochondrial

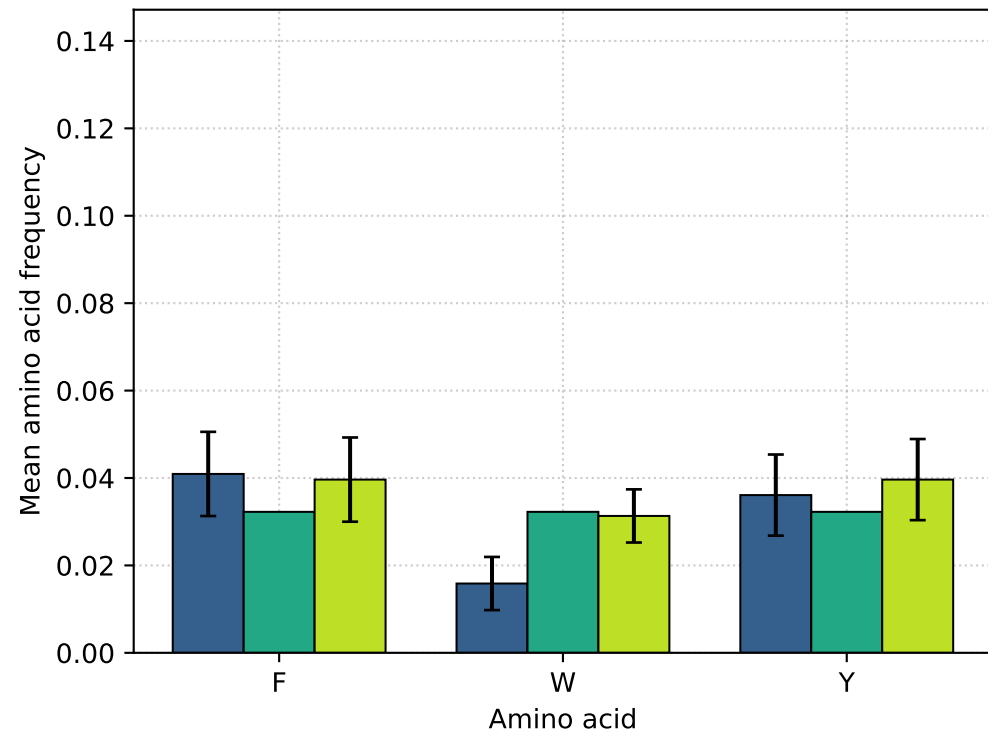
Aliphatic amino acids



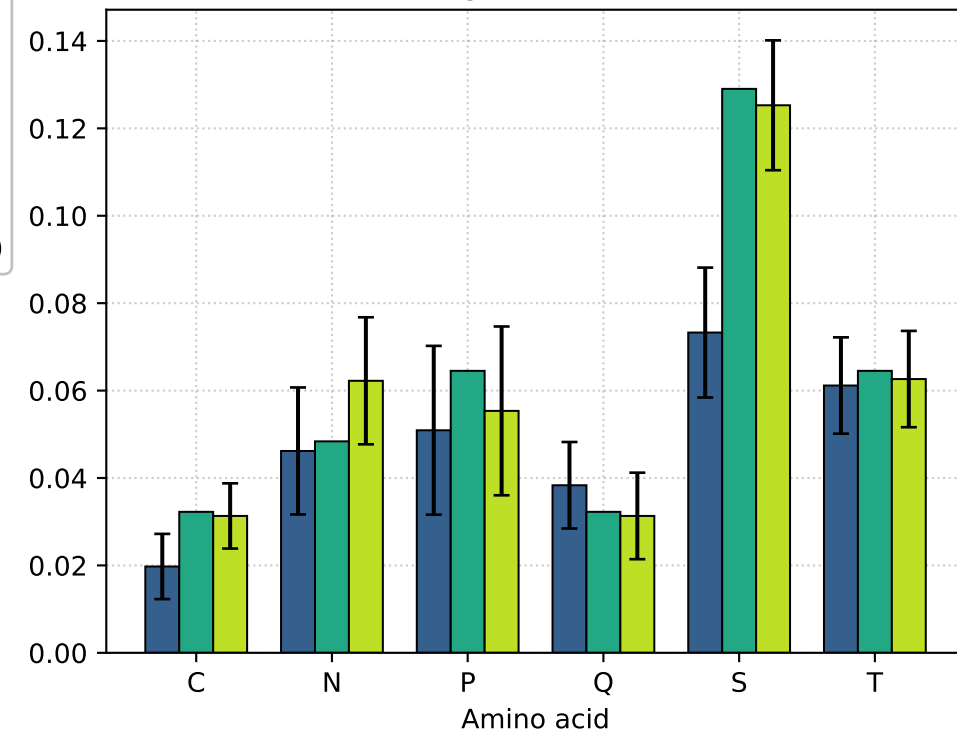
Charged amino acids



Aromatic amino acids



Uncharged amino acids



Observed
Codon
Codon+GC

Codon correlation:

- Mean Pearson:
- Coefficient: 0.55235
- p-value: 6.75789e-02

- Mean Spearman:
- Coefficient: 0.52641
- p-value: 9.42783e-02

- Mean log-MSE: 0.59512

Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.58706
- p-value: 3.66845e-02

- Mean Spearman:
- Coefficient: 0.52607
- p-value: 6.71734e-02

- Mean log-MSE: 0.57280