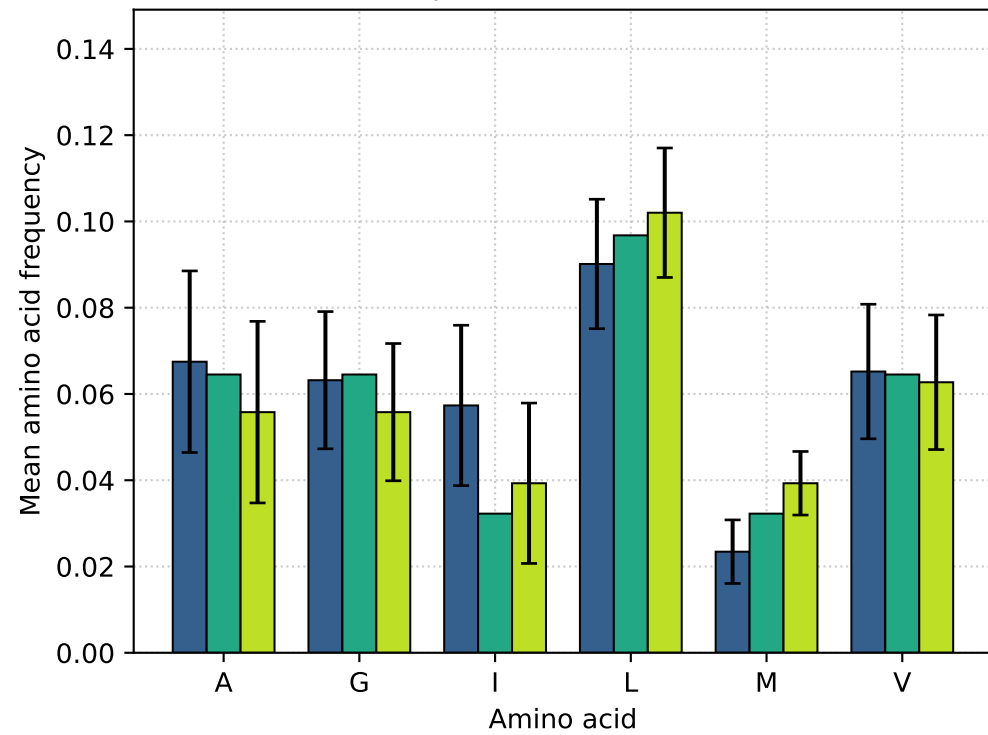


Viruses - Mean proteome amino acid frequency for genetic code: Trematode mitochondrial

Aliphatic amino acids

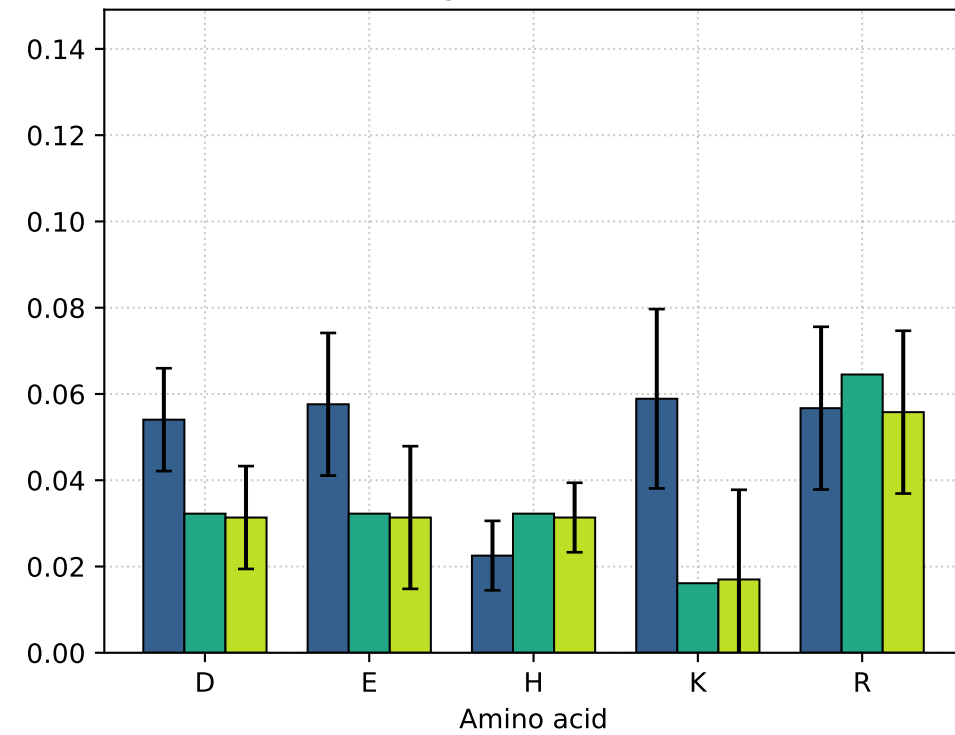


Observed
Codon
Codon+GC

Codon correlation:
- Mean Pearson:
- Coefficient: 0.54719
- p-value: 7.19221e-02

- Mean Spearman:
- Coefficient: 0.52855
- p-value: 8.91933e-02

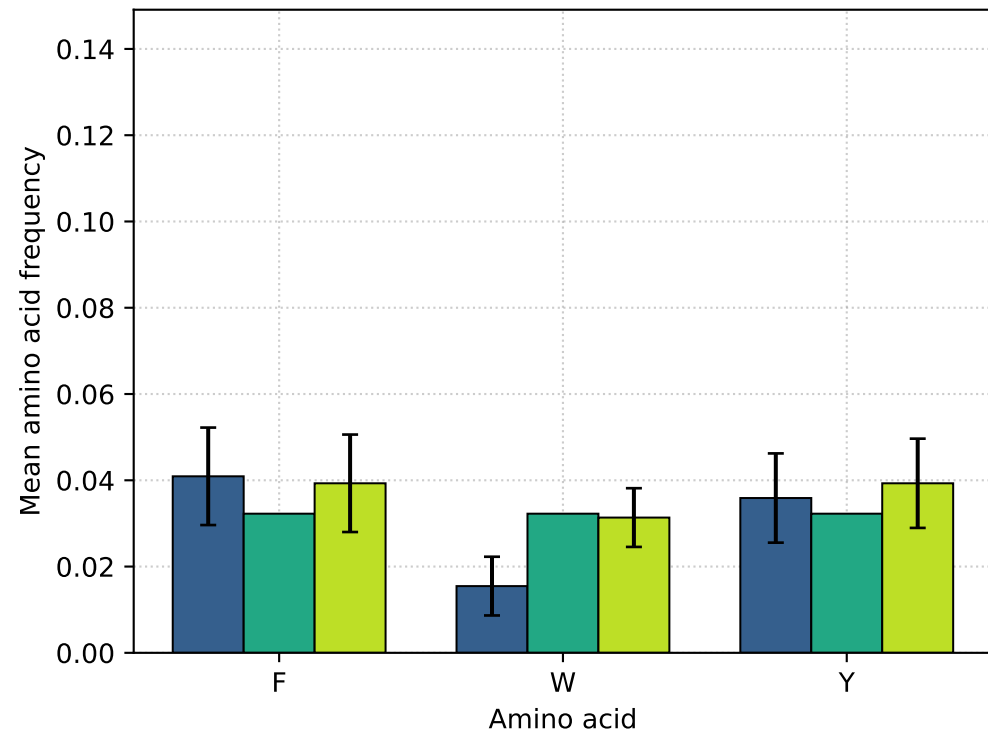
Charged amino acids



Codon + GC correlation:
- Mean Pearson:
- Coefficient: 0.57198
- p-value: 4.87677e-02

- Mean Spearman:
- Coefficient: 0.51684
- p-value: 7.62002e-02

Aromatic amino acids



Uncharged amino acids

