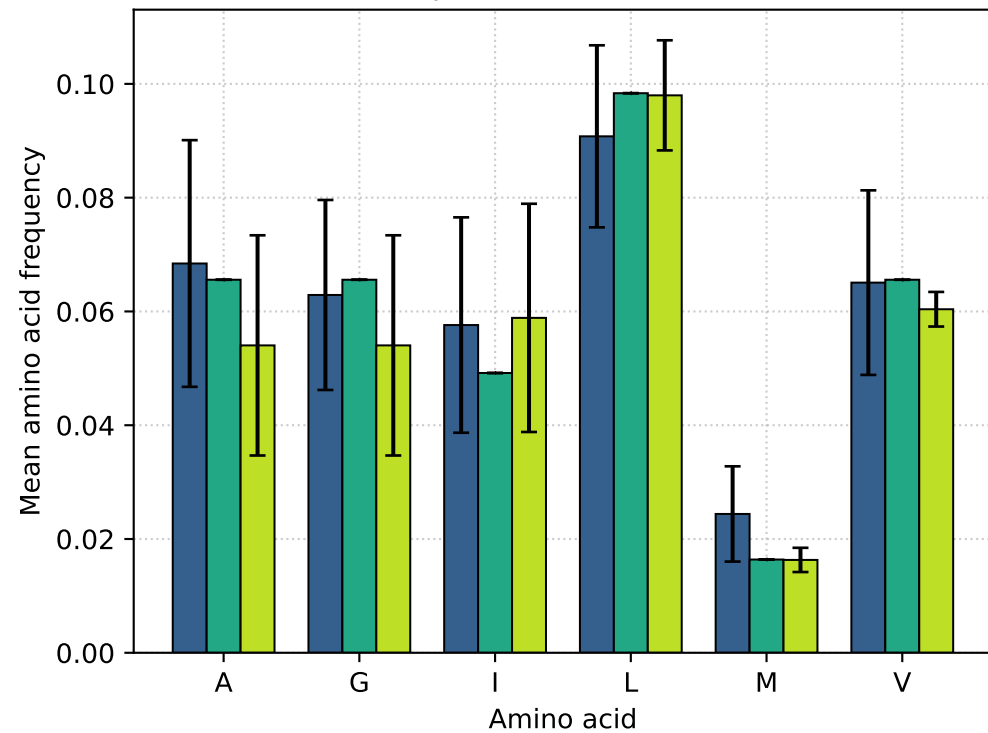


Viruses - Mean proteome amino acid frequency for genetic code: Standard

Aliphatic amino acids



Observed
Codon
Codon+GC

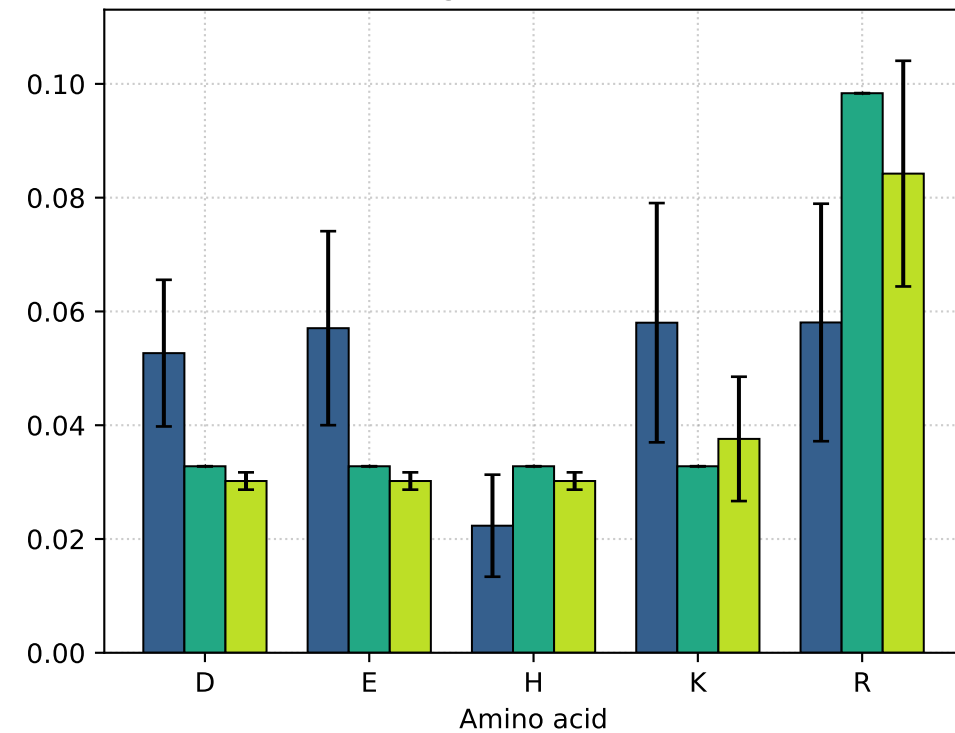
Codon correlation:
- Mean Pearson:
- Coefficient: 0.64392
- p-value: 3.39964e-02

- Mean Spearman:
- Coefficient: 0.66198
- p-value: 2.56652e-02

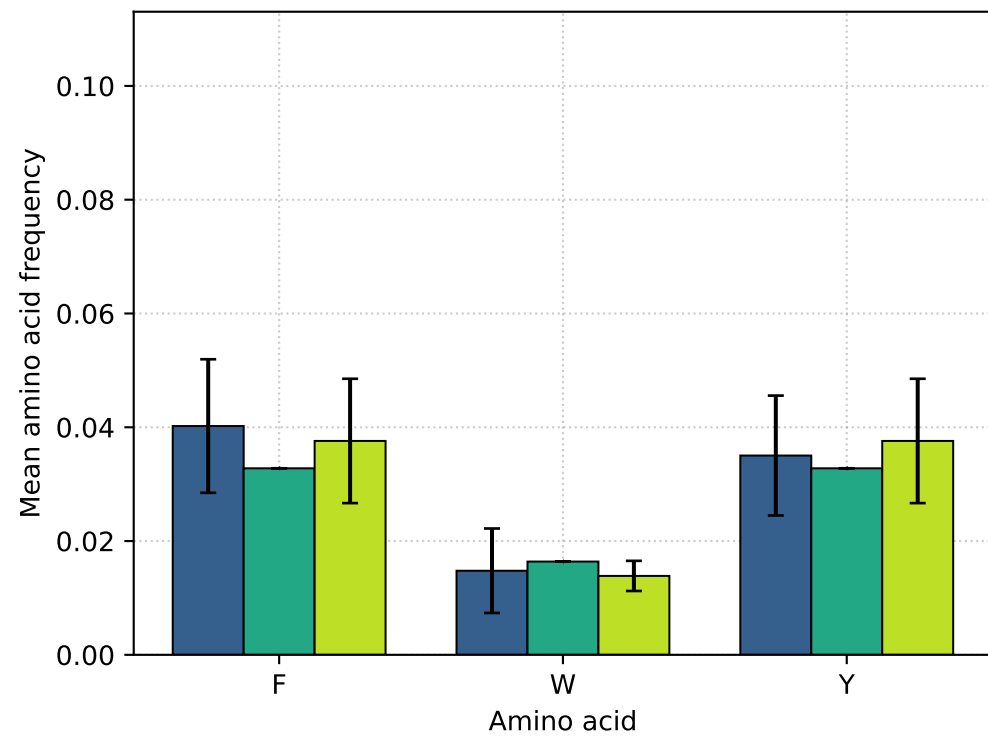
Codon + GC correlation:
- Mean Pearson:
- Coefficient: 0.71566
- p-value: 7.92674e-03

- Mean Spearman:
- Coefficient: 0.69433
- p-value: 8.59773e-03

Charged amino acids



Aromatic amino acids



Uncharged amino acids

