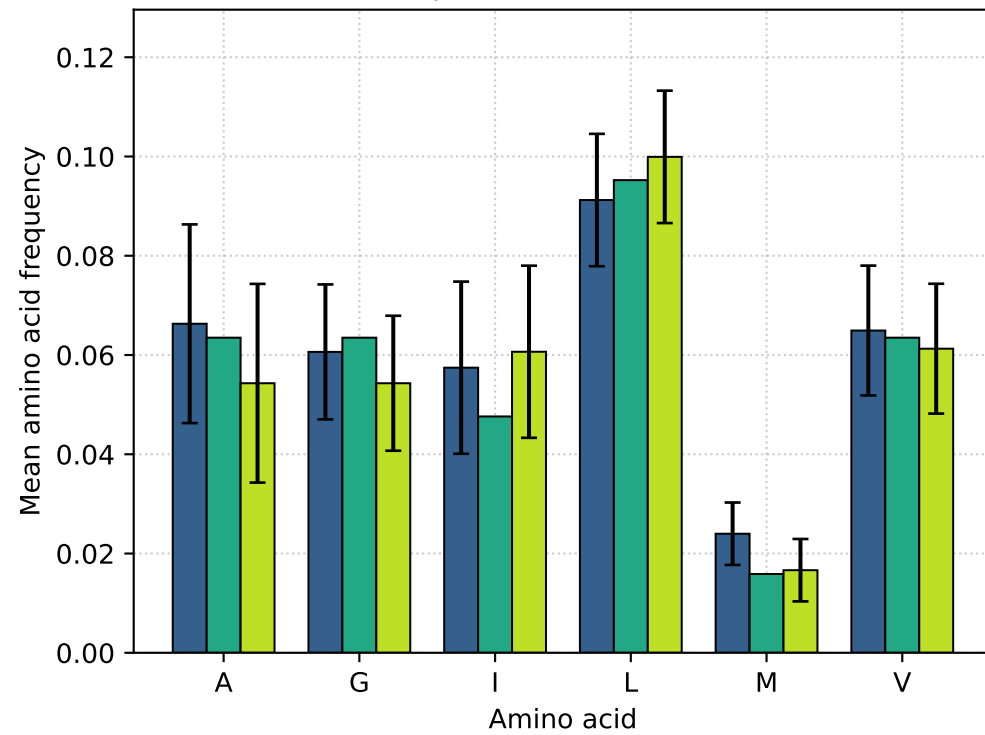
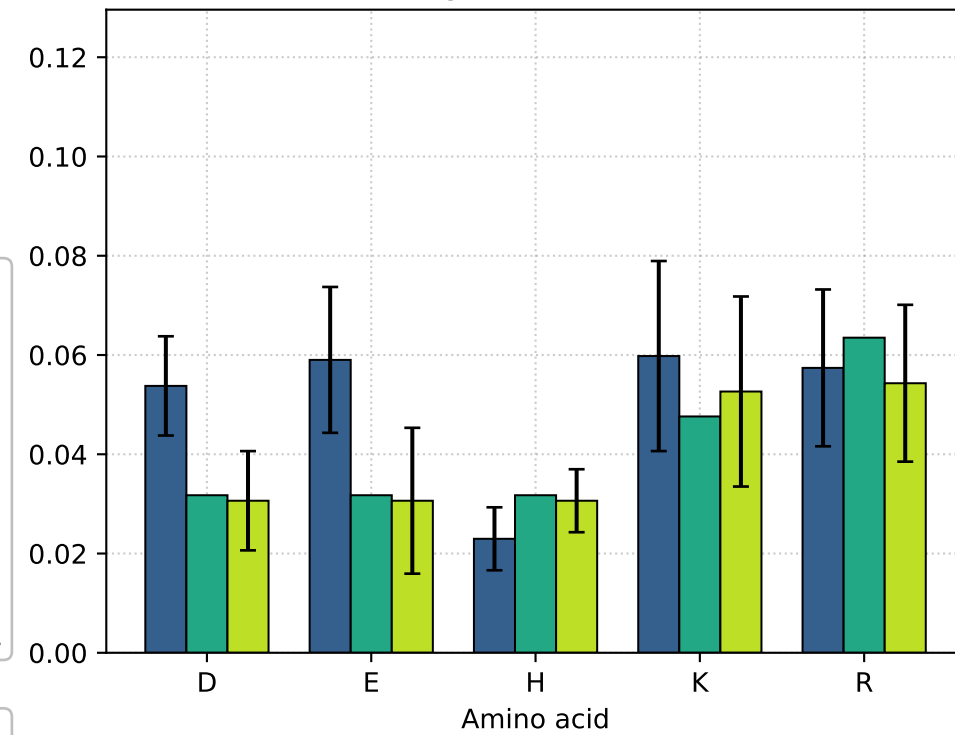


Viruses - Mean proteome amino acid frequency for genetic code: Cephalodiscidae mitochondrial uaa-tyr

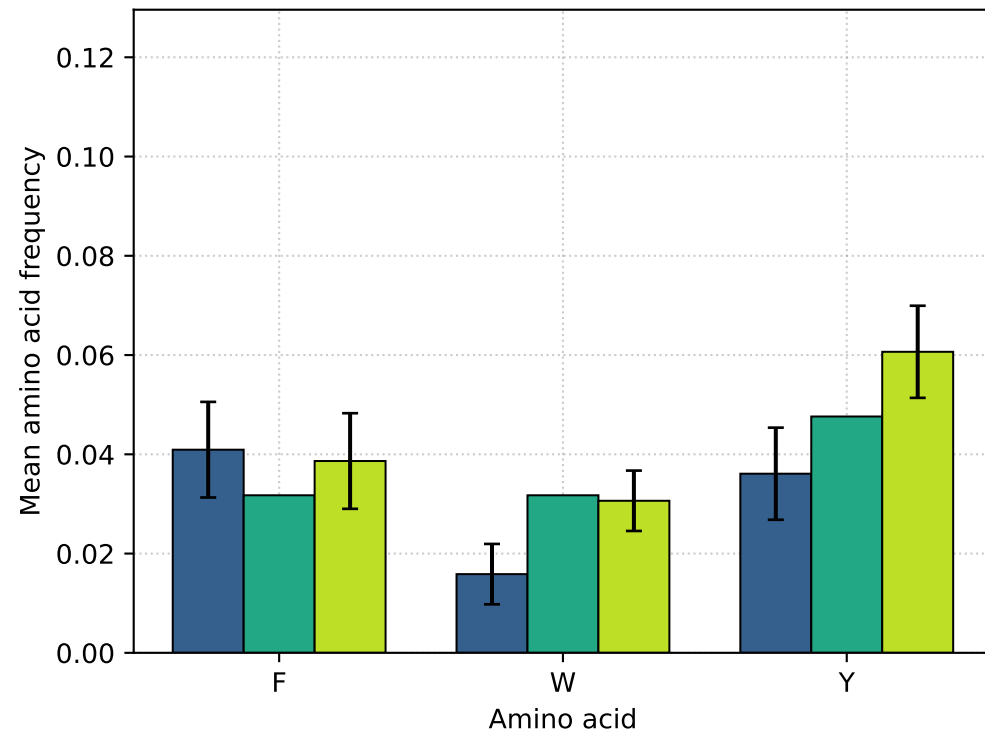
Aliphatic amino acids



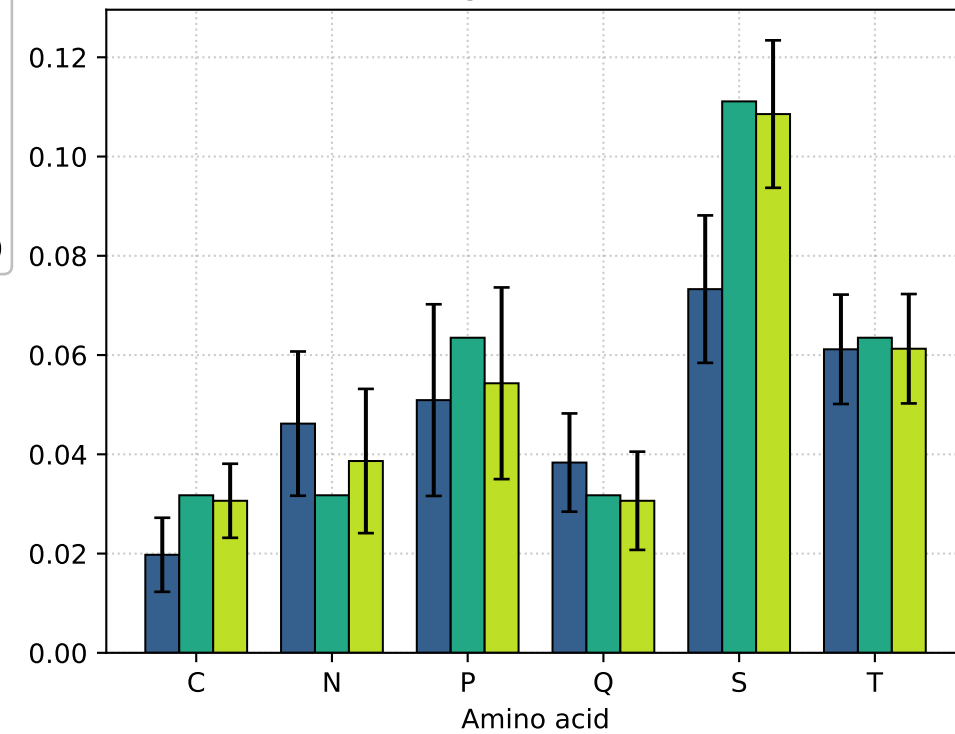
Charged amino acids



Aromatic amino acids



Uncharged amino acids



Codon correlation:

- Mean Pearson:
- Coefficient: 0.65240
- p-value: 1.92053e-02
- Mean Spearman:
- Coefficient: 0.65160
- p-value: 1.91865e-02
- Mean log-MSE: 0.41071

Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.69083
- p-value: 6.96222e-03
- Mean Spearman:
- Coefficient: 0.67134
- p-value: 7.62892e-03
- Mean log-MSE: 0.39290