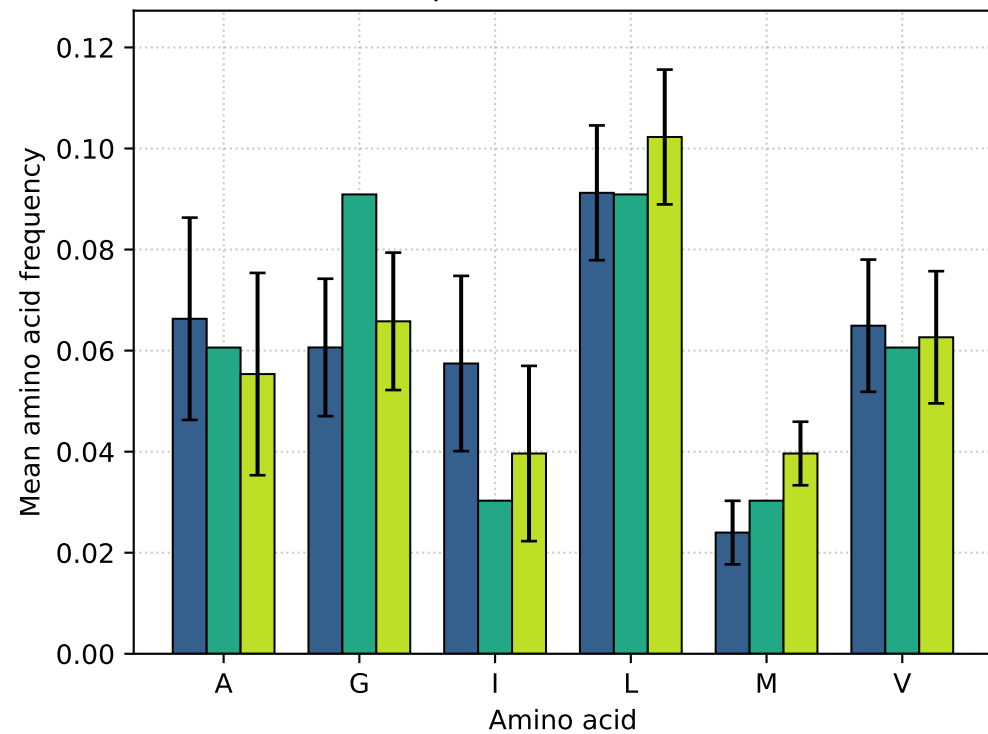
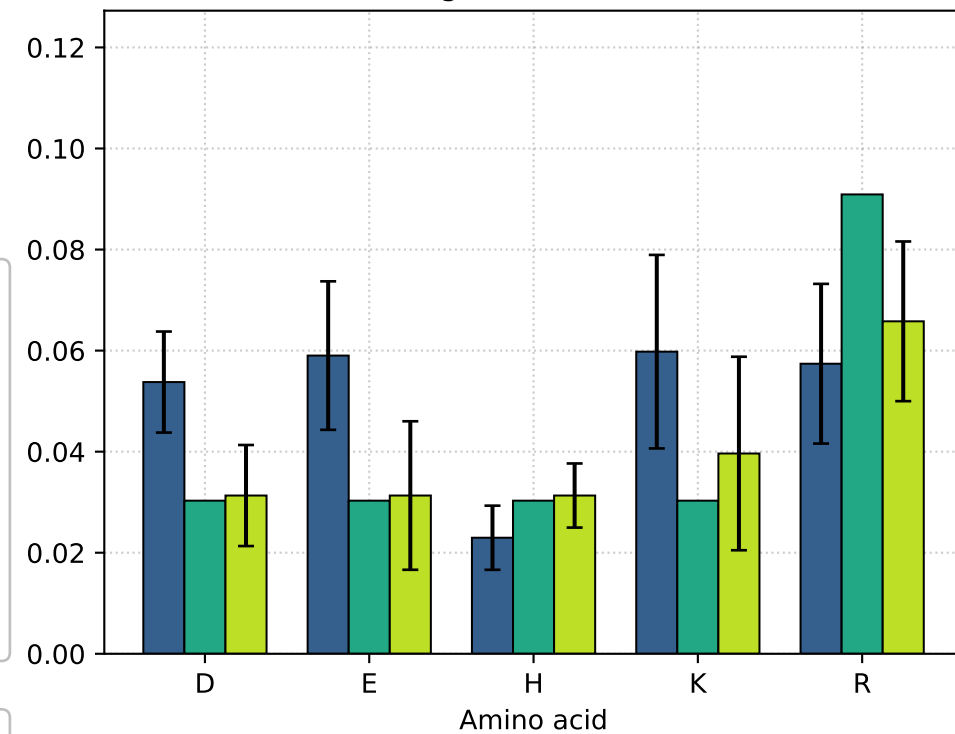


Viruses - Mean proteome amino acid frequency for genetic code: Ascidian mitochondrial

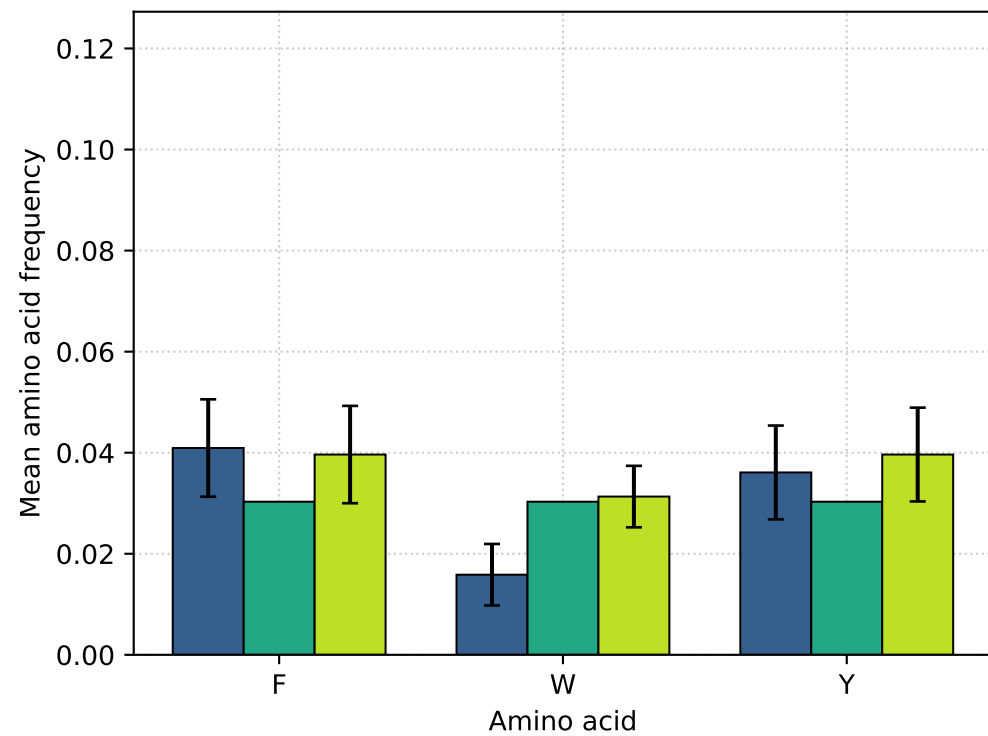
Aliphatic amino acids



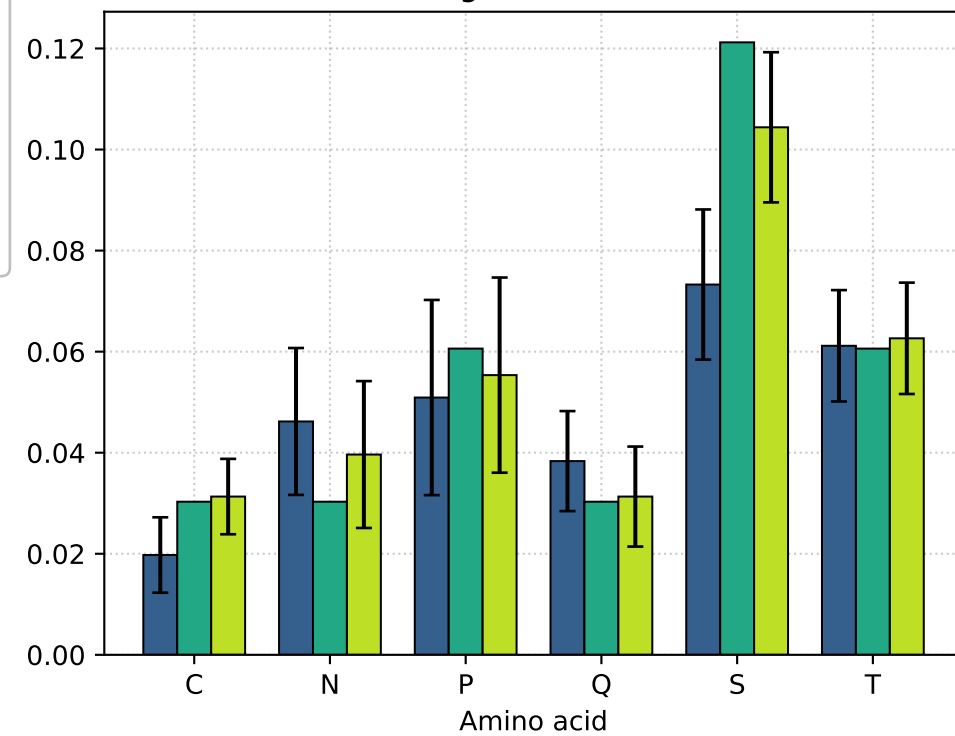
Charged amino acids



Aromatic amino acids



Uncharged amino acids



Codon correlation:

- Mean Pearson:
- Coefficient: 0.55820
- p-value: 6.77658e-02
- Mean Spearman:
- Coefficient: 0.57233
- p-value: 7.63304e-02
- Mean log-MSE: 0.52023

Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.67573
- p-value: 8.97033e-03
- Mean Spearman:
- Coefficient: 0.62661
- p-value: 1.35950e-02
- Mean log-MSE: 0.40846