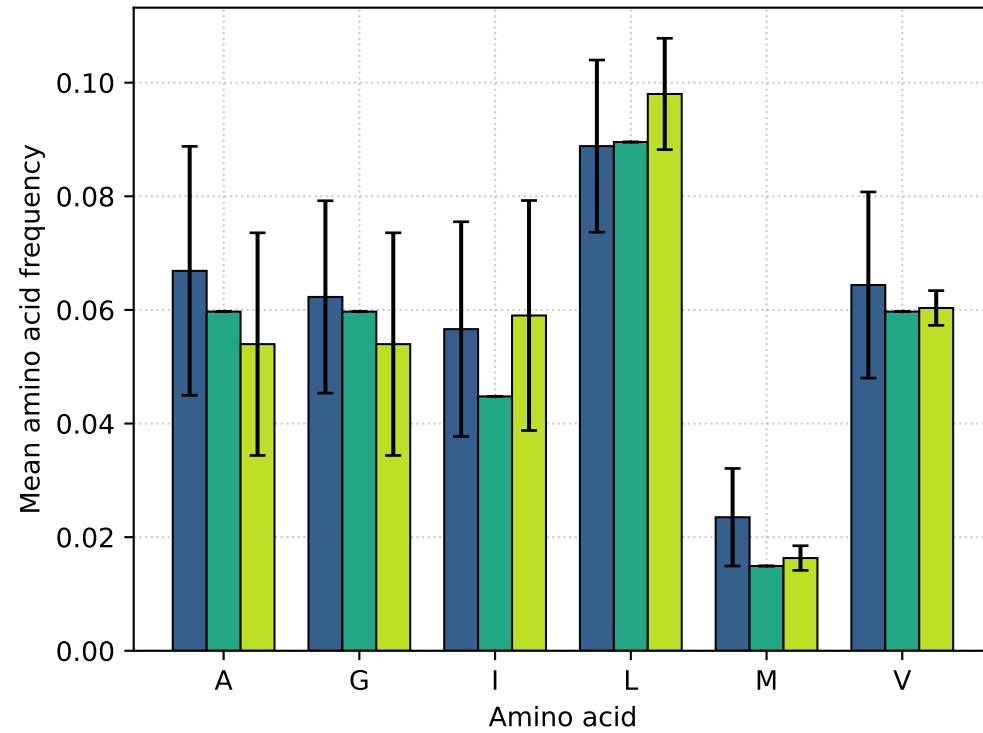


Viruses - Mean proteomic amino acid frequency for genetic code: Condyllostoma nuclear

Aliphatic amino acids



Observed
Codon
Codon+GC

Codon correlation:

- Pearson:
- Coefficient: 0.577951
- p-value: 0.057952

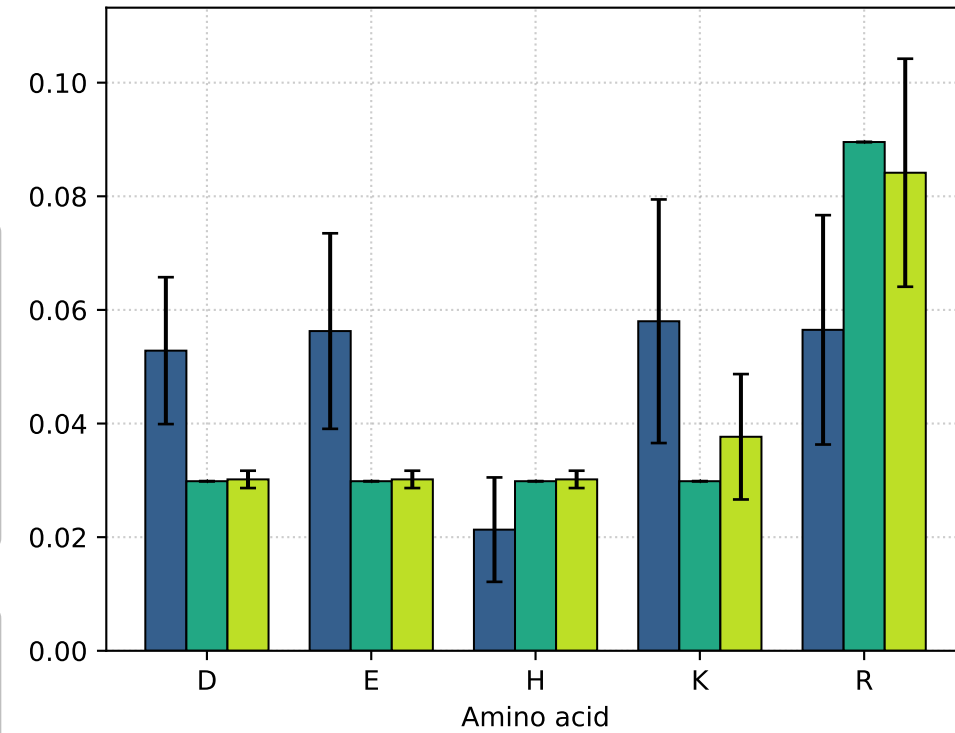
- Spearman:
- Coefficient: 0.573059
- p-value: 0.060688

Codon + GC correlation:

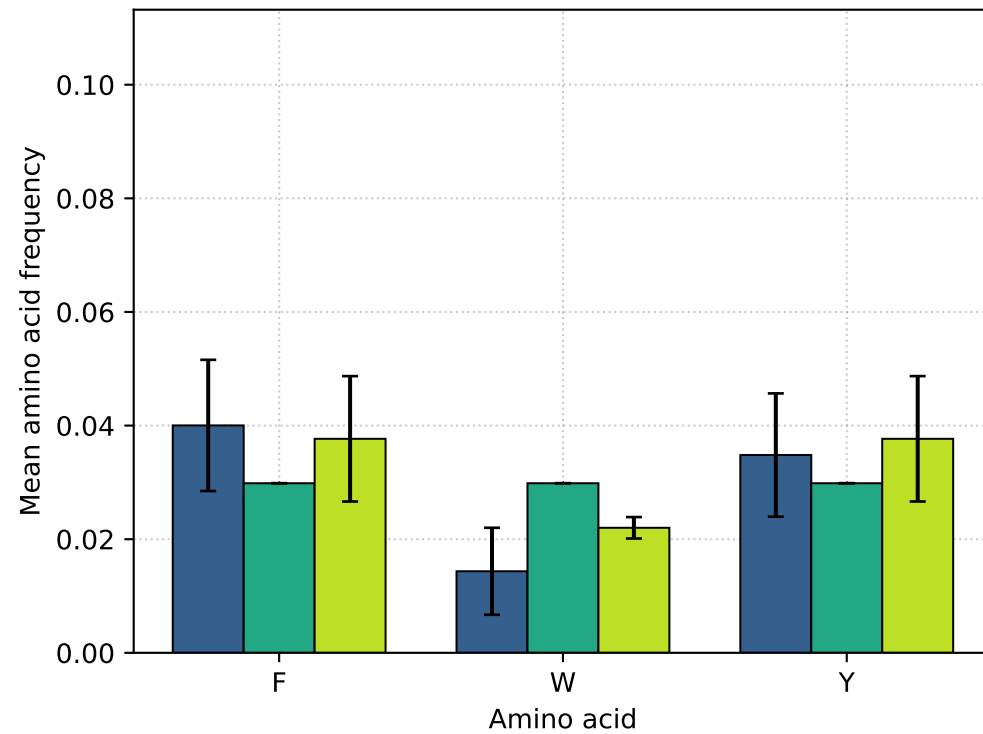
- Pearson:
- Coefficient: 0.683058
- p-value: 0.010843

- Spearman:
- Coefficient: 0.658365
- p-value: 0.012920

Charged amino acids



Aromatic amino acids



Uncharged amino acids

