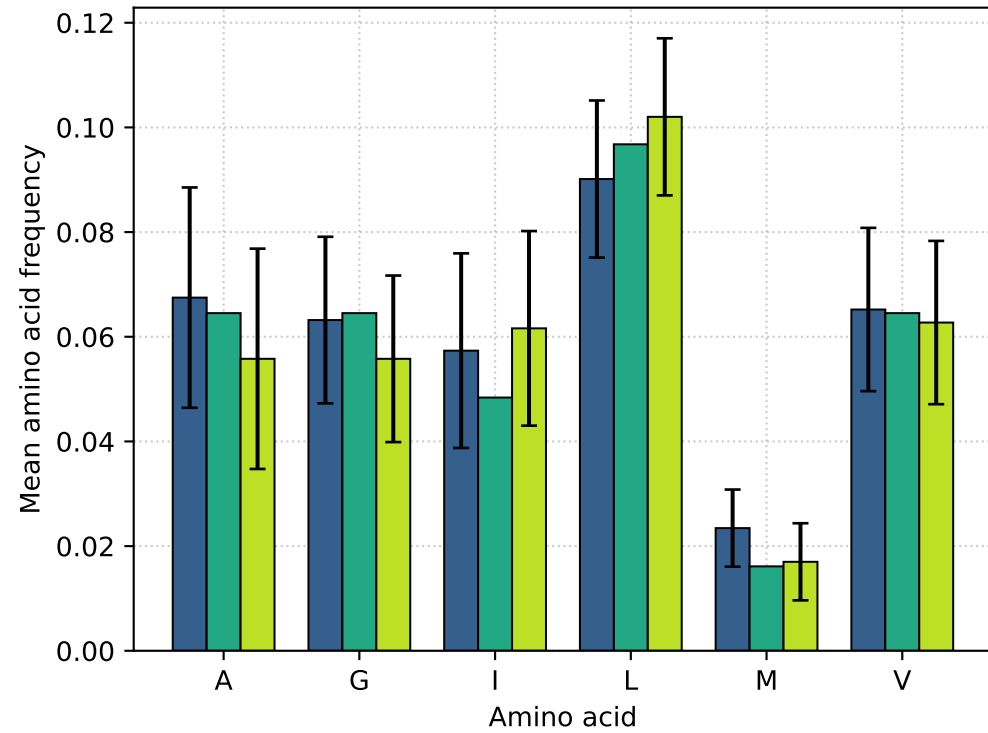


Viruses - Mean proteome amino acid frequency for genetic code: Mold-protozoan-coelenterate mitochondrial and mycoplasma spiroplasma

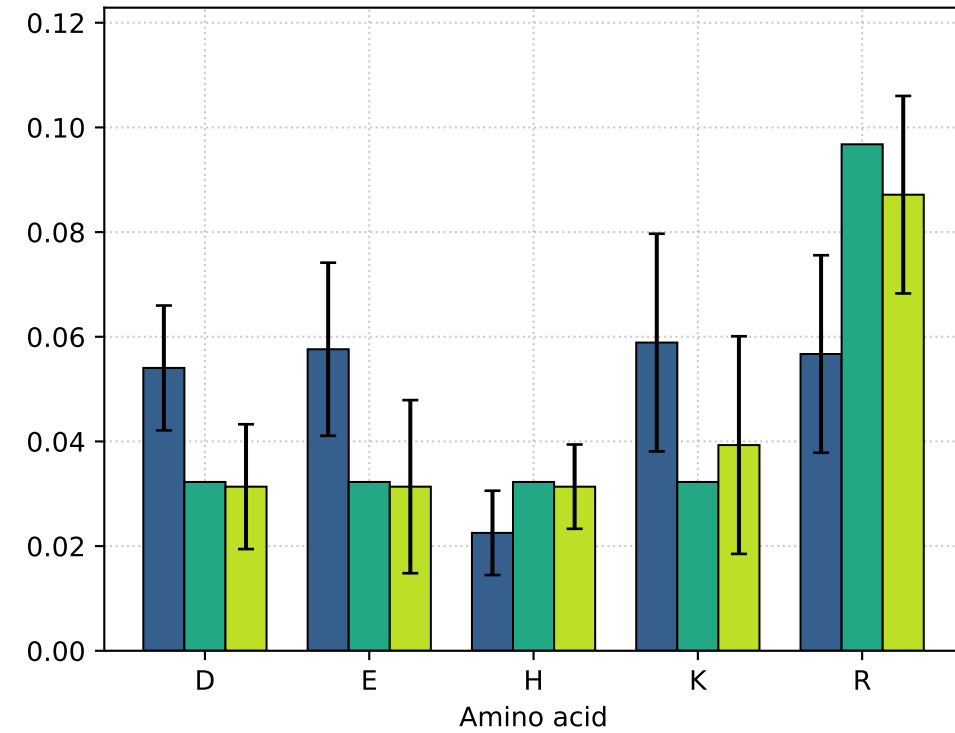
Aliphatic amino acids



Codon correlation:
- Mean Pearson:
- Coefficient: 0.61050
- p-value: 4.37991e-02

- Mean Spearman:
- Coefficient: 0.62252
- p-value: 3.76680e-02

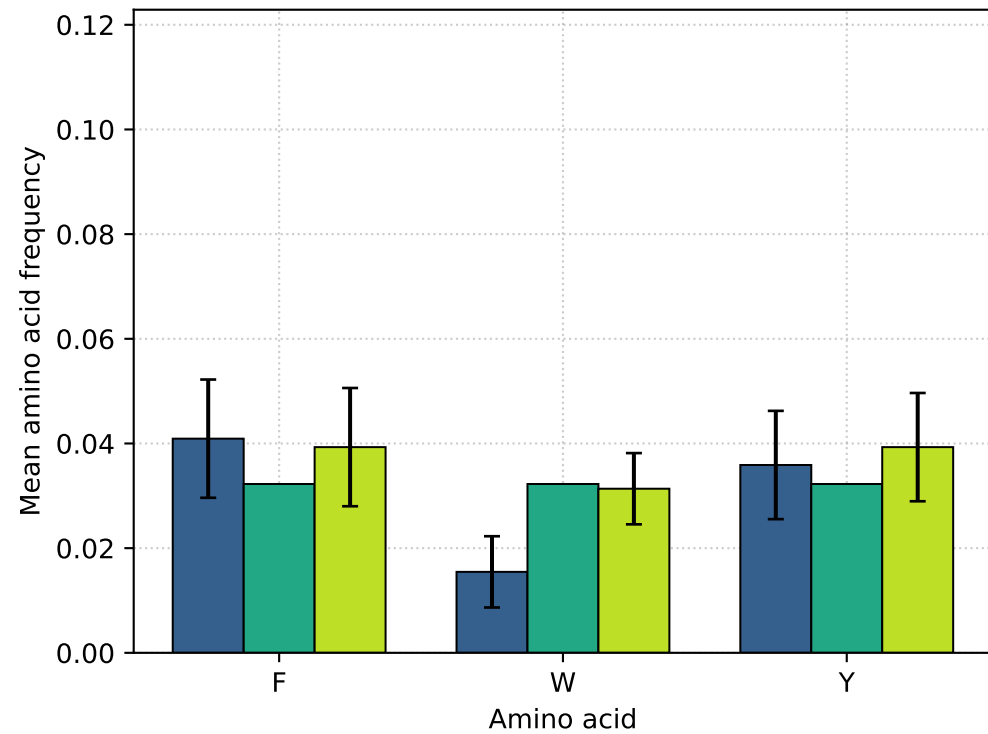
Charged amino acids



Codon + GC correlation:
- Mean Pearson:
- Coefficient: 0.68237
- p-value: 1.08792e-02

- Mean Spearman:
- Coefficient: 0.66165
- p-value: 1.12080e-02

Aromatic amino acids



Uncharged amino acids

