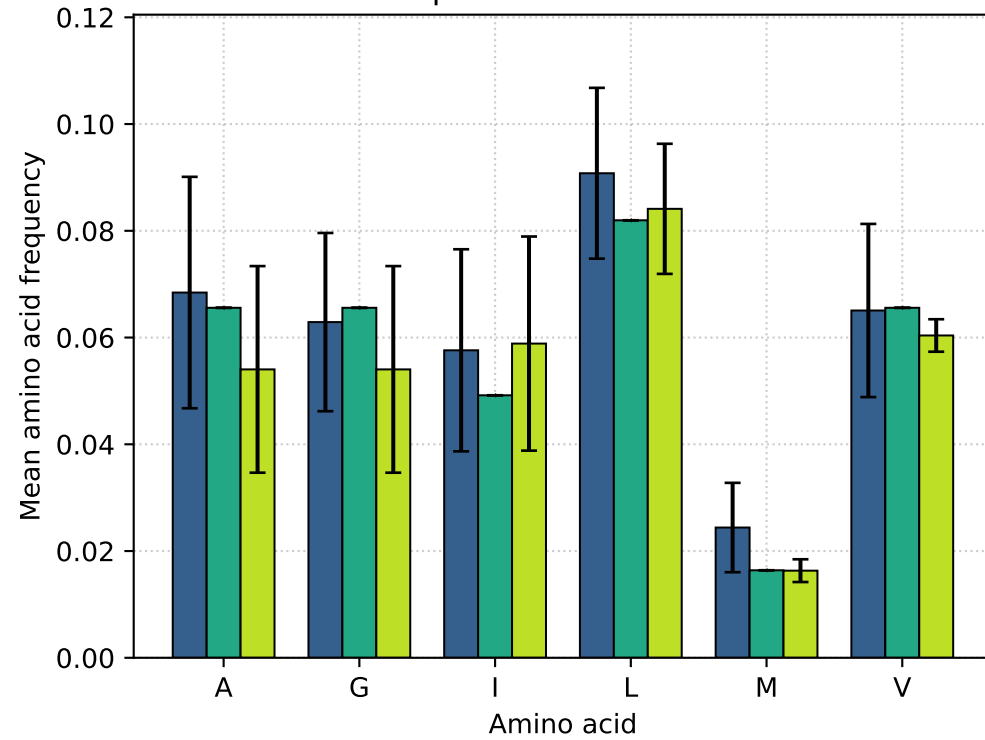


Viruses - Mean proteome amino acid frequency for genetic code: Alternative yeast

Aliphatic amino acids



Observed
Codon
Codon+GC

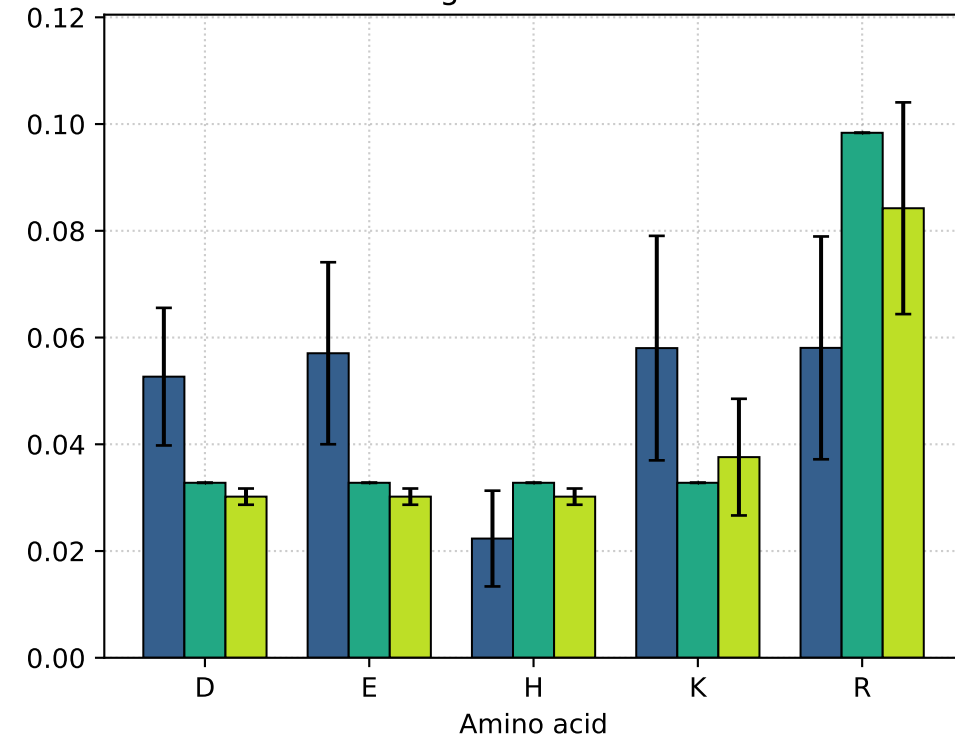
Codon correlation:
- Mean Pearson:
- Coefficient: 0.61023
- p-value: 4.31398e-02

- Mean Spearman:
- Coefficient: 0.65728
- p-value: 2.69981e-02

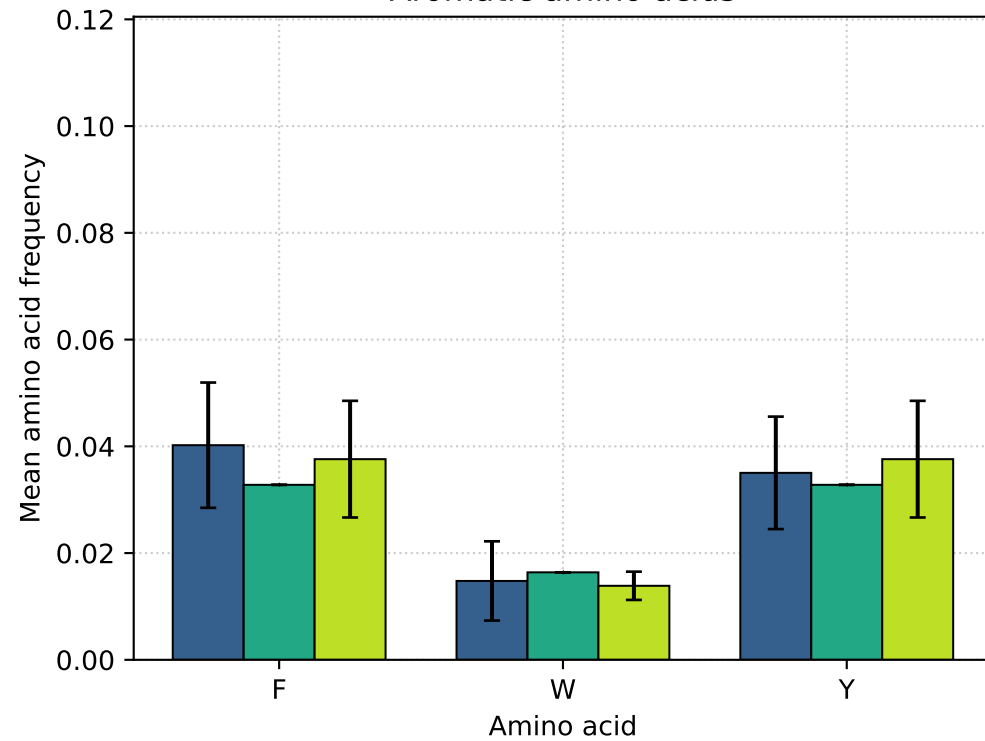
Codon + GC correlation:
- Mean Pearson:
- Coefficient: 0.69137
- p-value: 1.04178e-02

- Mean Spearman:
- Coefficient: 0.69024
- p-value: 9.04509e-03

Charged amino acids



Aromatic amino acids



Uncharged amino acids

