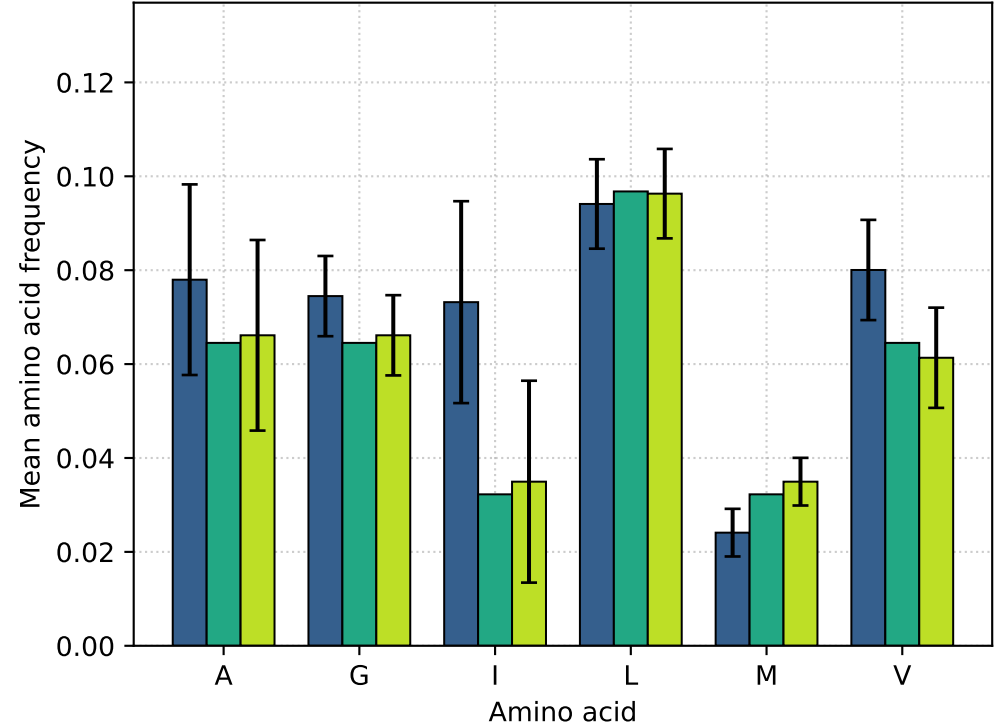


Archaea - Mean proteome amino acid frequency for genetic code: Trematode mitochondrial

Aliphatic amino acids



Observed
Codon
Codon+GC

Codon correlation:

- Mean Pearson:
- Coefficient: 0.41092
- p-value: 1.29383e-01

- Mean Spearman:
- Coefficient: 0.44434
- p-value: 1.17167e-01

- Mean log-MSE: 0.88487

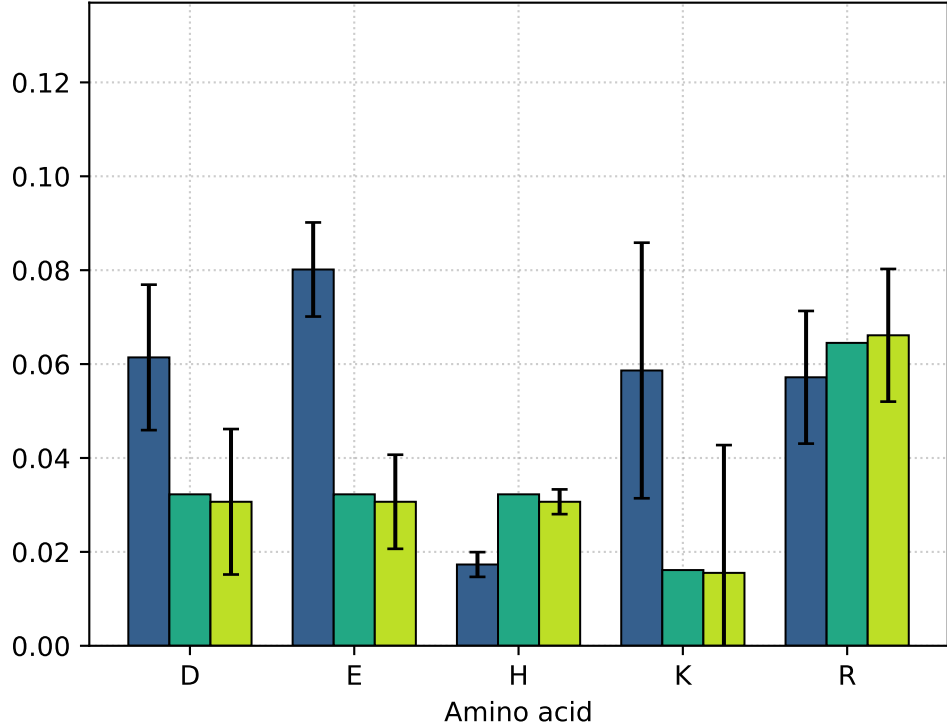
Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.45986
- p-value: 8.61822e-02

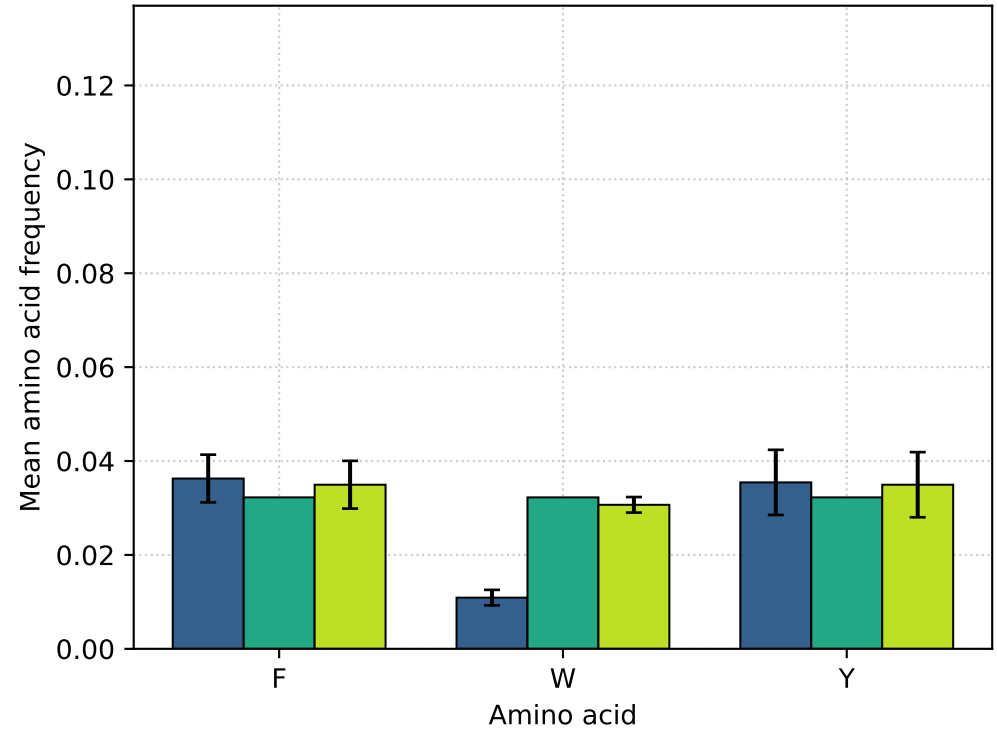
- Mean Spearman:
- Coefficient: 0.42281
- p-value: 1.30417e-01

- Mean log-MSE: 0.91511

Charged amino acids



Aromatic amino acids



Uncharged amino acids

