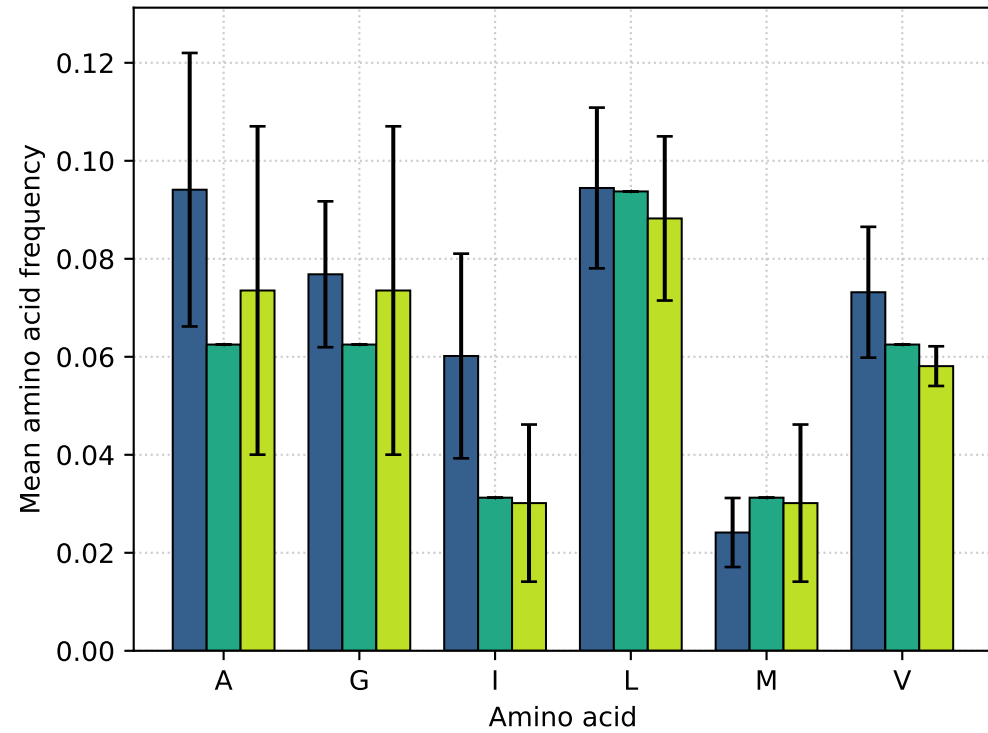


Bacteria - Mean proteomic amino acid frequency for genetic code: Trematode mitochondrial

Aliphatic amino acids



Observed
Codon
Codon+GC

Codon correlation:

- Pearson:
- Coefficient: 0.429558
- p-value: 0.115723

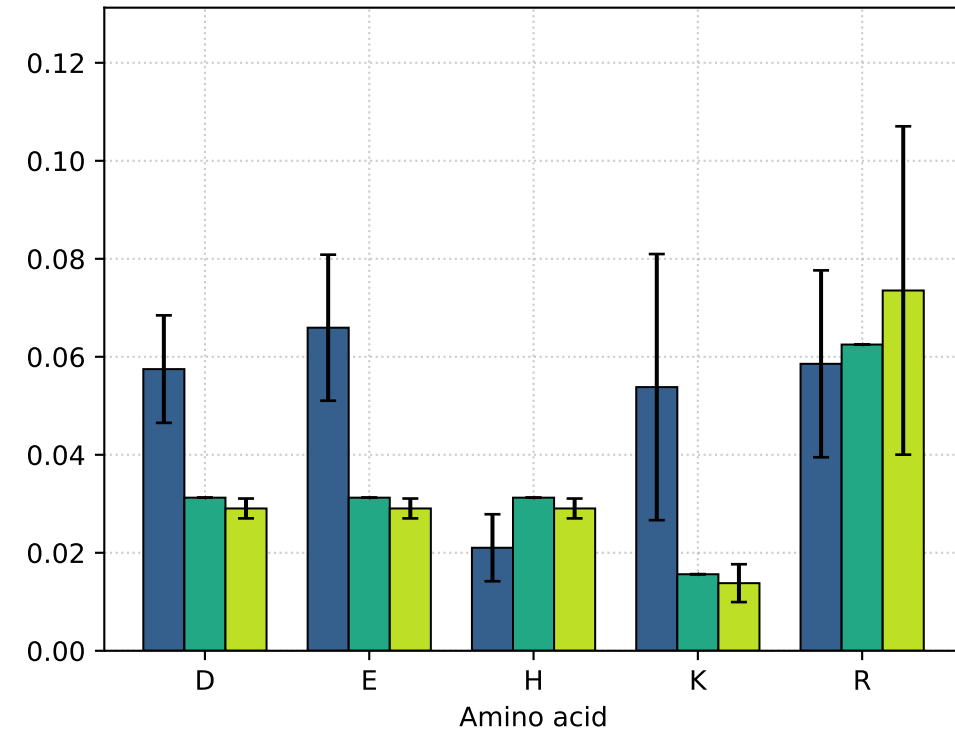
- Spearman:
- Coefficient: 0.485072
- p-value: 0.103557

Codon + GC correlation:

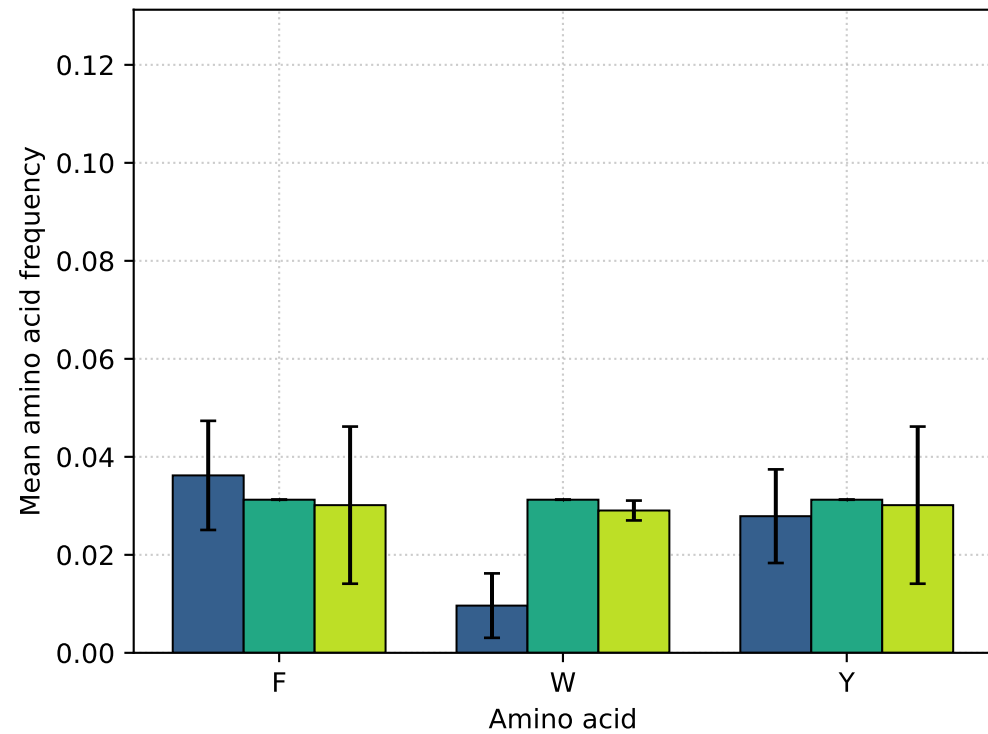
- Pearson:
- Coefficient: 0.497885
- p-value: 0.088417

- Spearman:
- Coefficient: 0.467521
- p-value: 0.123032

Charged amino acids



Aromatic amino acids



Uncharged amino acids

