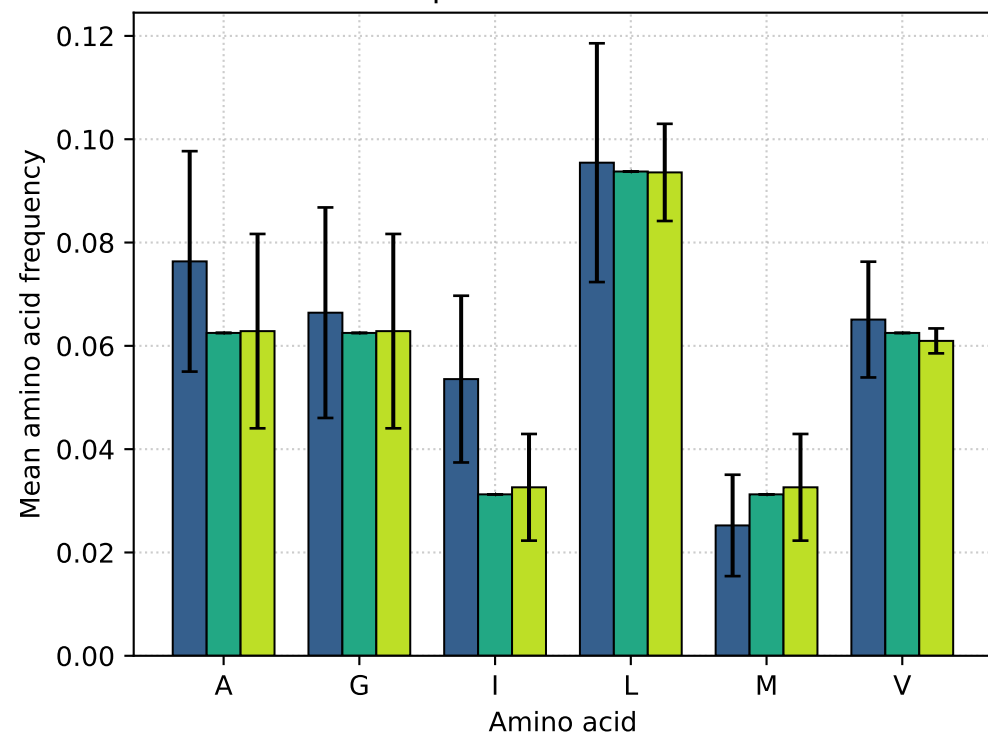


Eukaryota - Mean proteomic amino acid frequency for genetic code: Trematode mitochondrial

Aliphatic amino acids



Observed
Codon
Codon+GC

Codon correlation:

- Pearson:
- Coefficient: 0.568417
- p-value: 0.047755

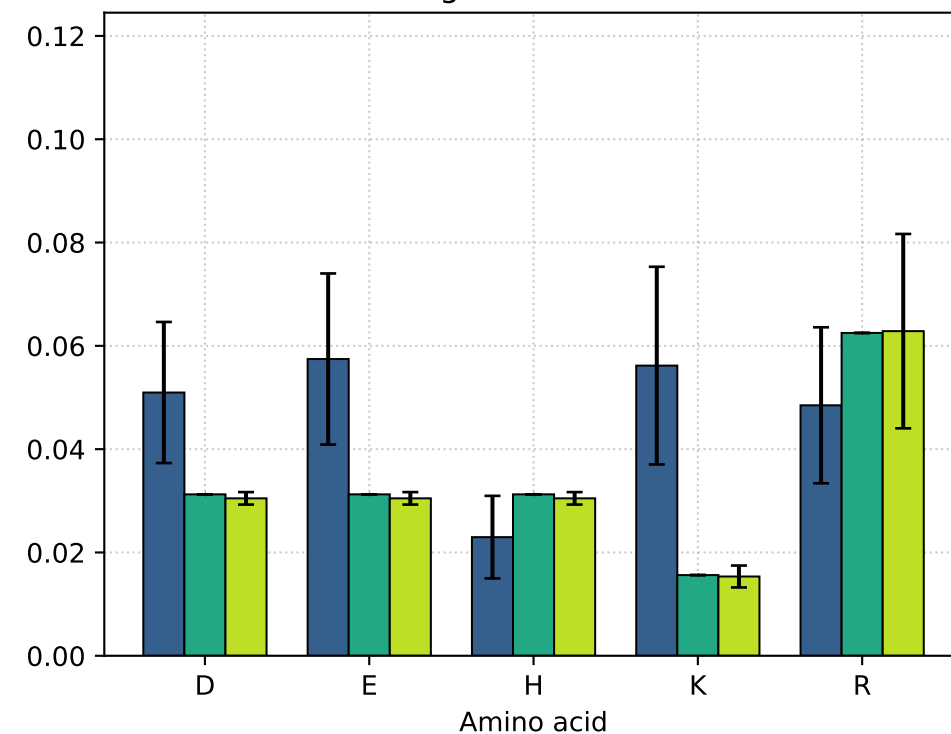
- Spearman:
- Coefficient: 0.551306
- p-value: 0.061191

Codon + GC correlation:

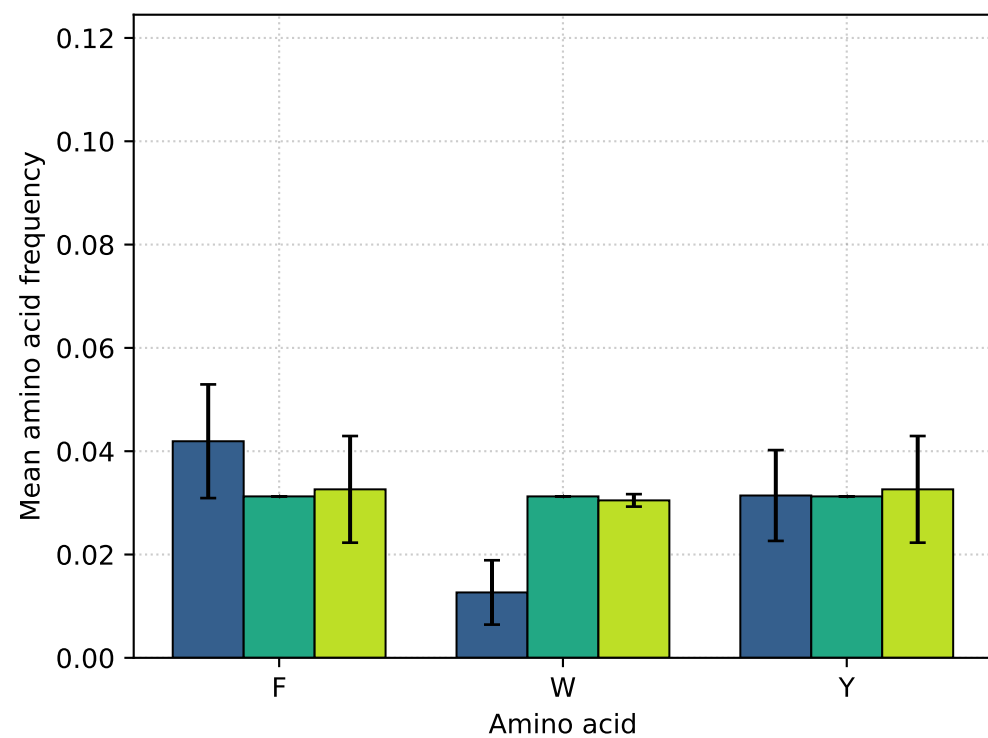
- Pearson:
- Coefficient: 0.587375
- p-value: 0.036132

- Spearman:
- Coefficient: 0.529846
- p-value: 0.065644

Charged amino acids



Aromatic amino acids



Uncharged amino acids

