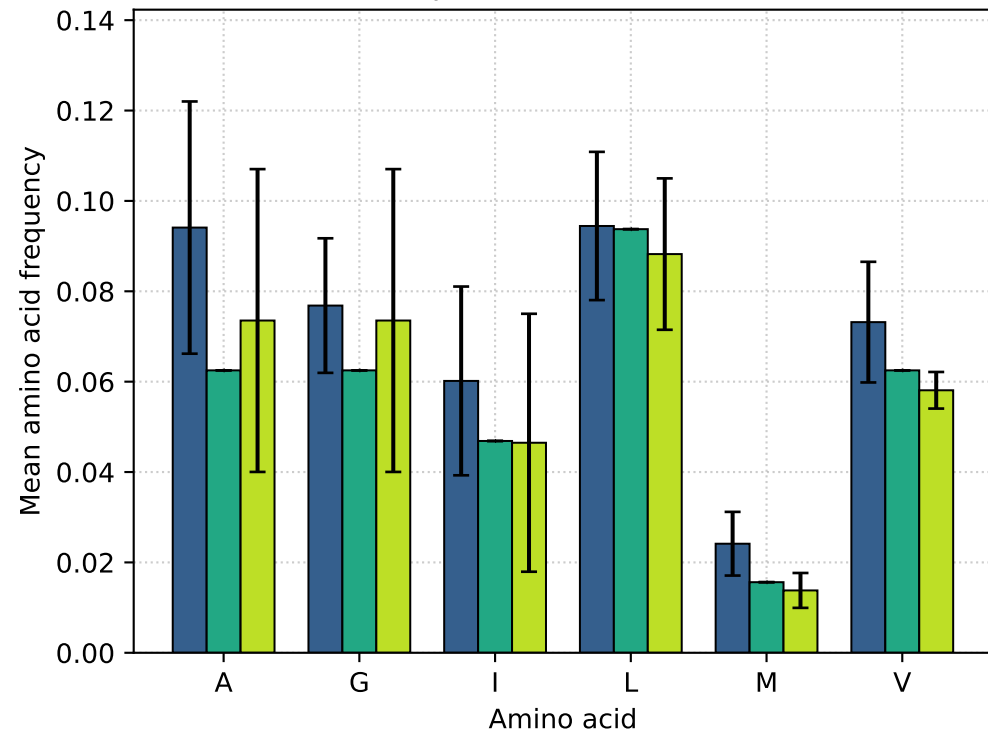


Bacteria - Mean proteomic amino acid frequency for genetic code: Mold-protozoan-coelenterate mitochondrial and mycoplasma spiroplasma

Aliphatic amino acids



Observed
Codon
Codon+GC

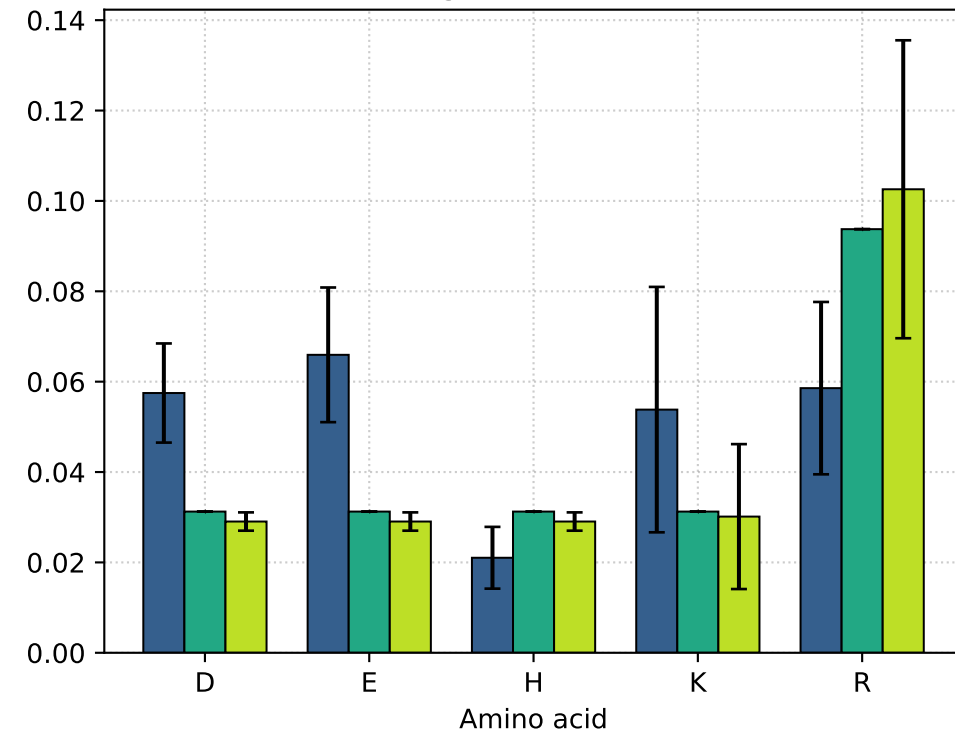
Codon correlation:

- Pearson:
 - Coefficient: 0.541336
 - p-value: 0.055474
- Spearman:
 - Coefficient: 0.591467
 - p-value: 0.038397

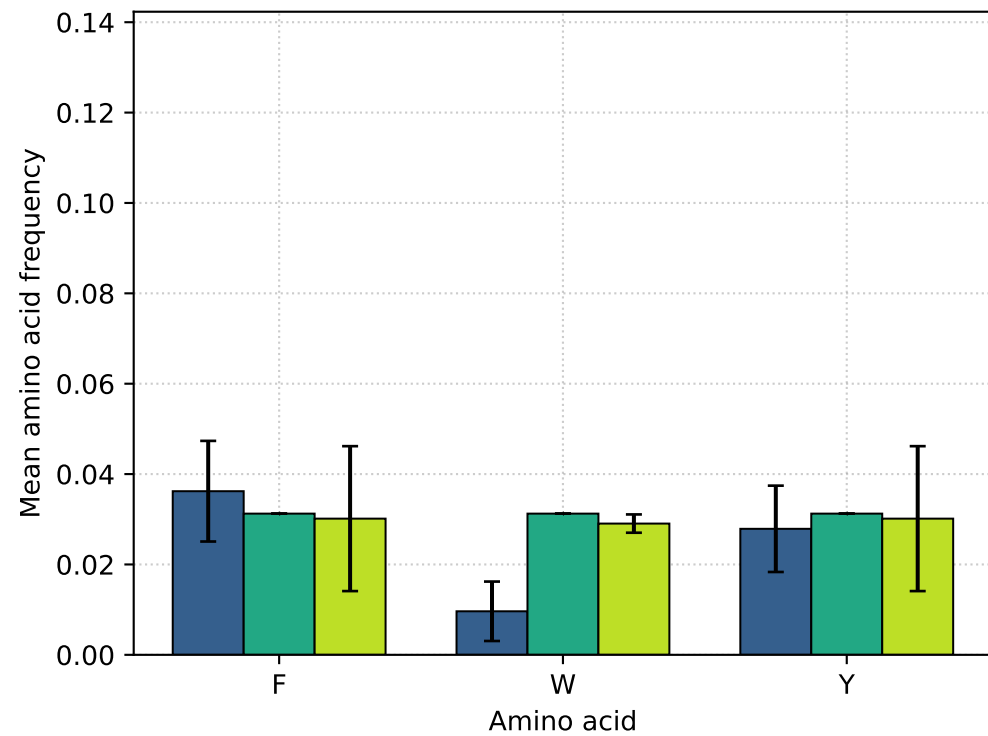
Codon + GC correlation:

- Pearson:
 - Coefficient: 0.607891
 - p-value: 0.020005
- Spearman:
 - Coefficient: 0.601290
 - p-value: 0.021645

Charged amino acids



Aromatic amino acids



Uncharged amino acids

