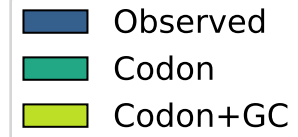
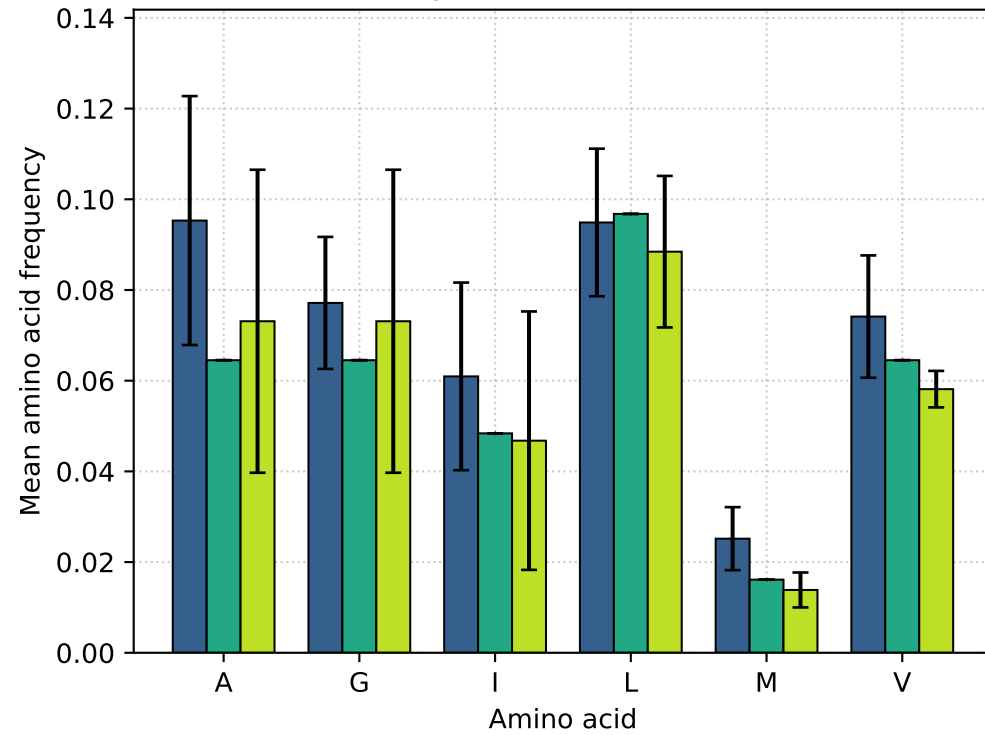


# Bacteria - Mean proteome amino acid frequency for genetic code: Mold-protozoan-coelenterate mitochondrial and mycoplasma spiroplasma

## Aliphatic amino acids



**Codon correlation:**

- Mean Pearson:
- Coefficient: 0.54266
- p-value: 5.43821e-02

- Mean Spearman:

- Coefficient: 0.58978
- p-value: 3.73787e-02

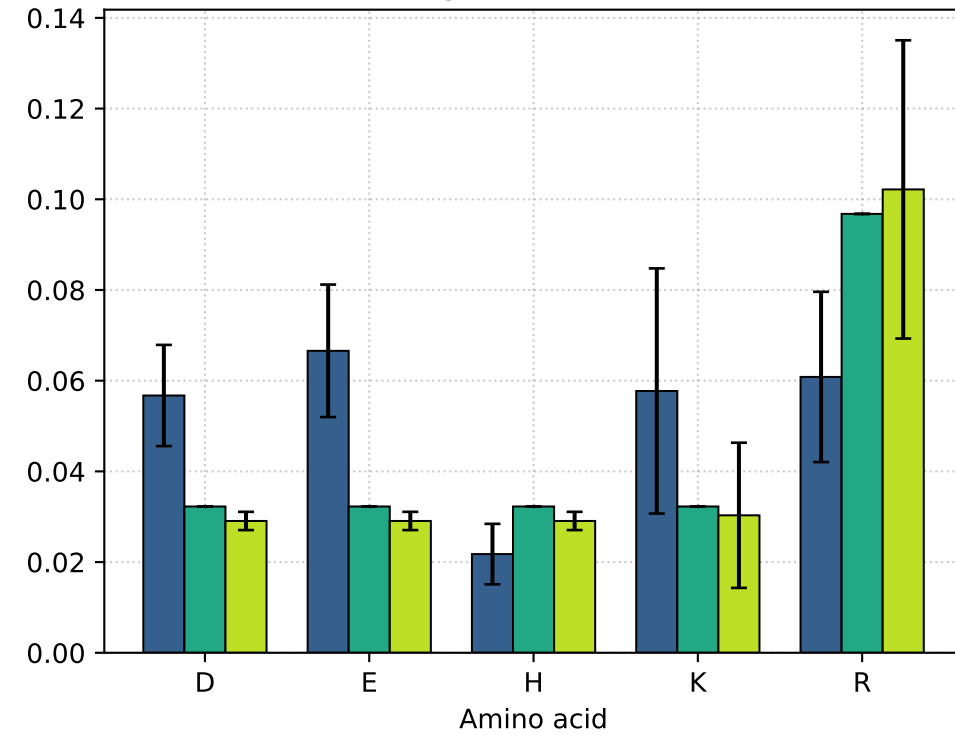
**Codon + GC correlation:**

- Mean Pearson:
- Coefficient: 0.60731
- p-value: 2.03140e-02

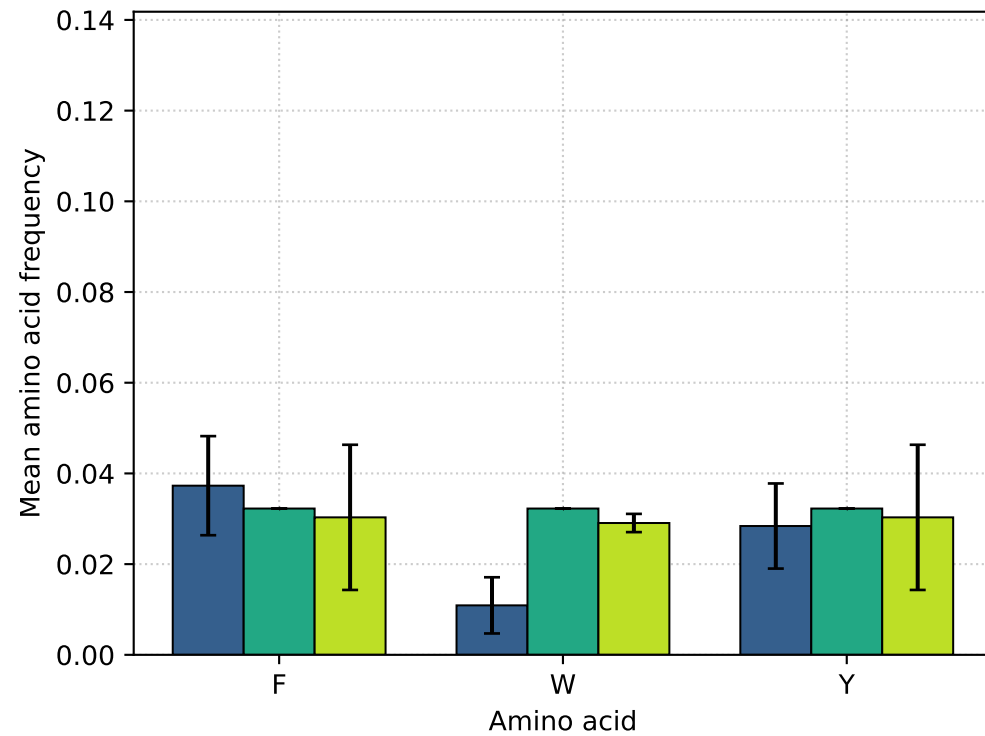
- Mean Spearman:

- Coefficient: 0.59683
- p-value: 2.27523e-02

## Charged amino acids



## Aromatic amino acids



## Uncharged amino acids

