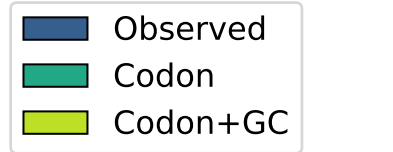
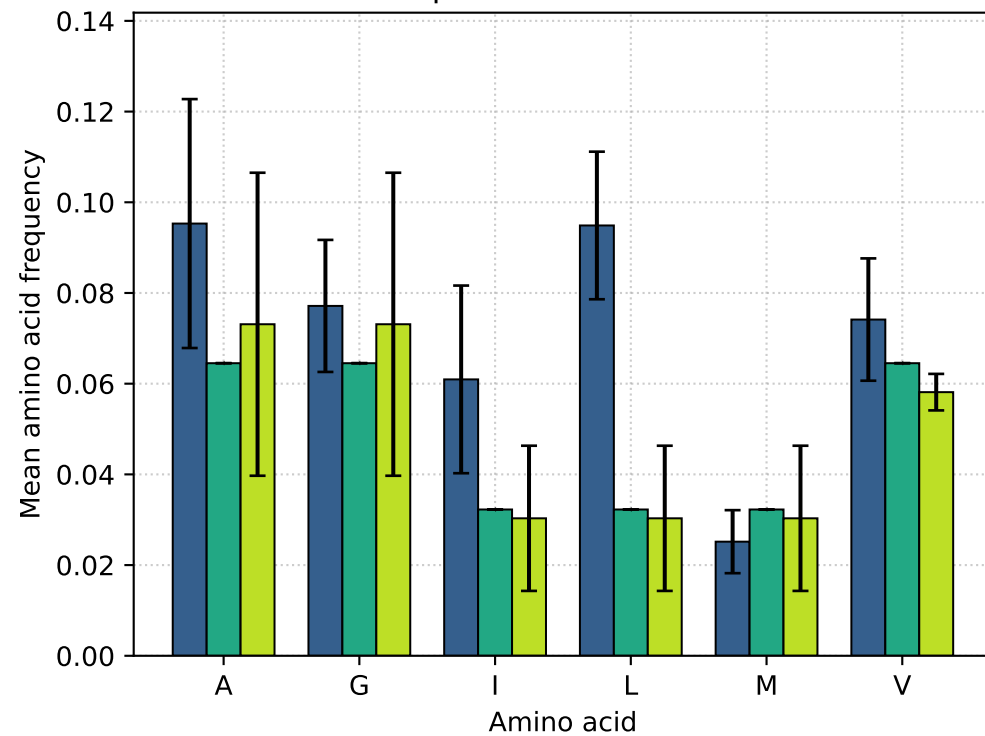


# Bacteria - Mean proteome amino acid frequency for genetic code: Yeast mitochondrial

## Aliphatic amino acids



**Codon correlation:**

- Mean Pearson:
- Coefficient: 0.25717
- p-value: 3.18392e-01

- Mean Spearman:

- Coefficient: 0.37659
- p-value: 1.85928e-01

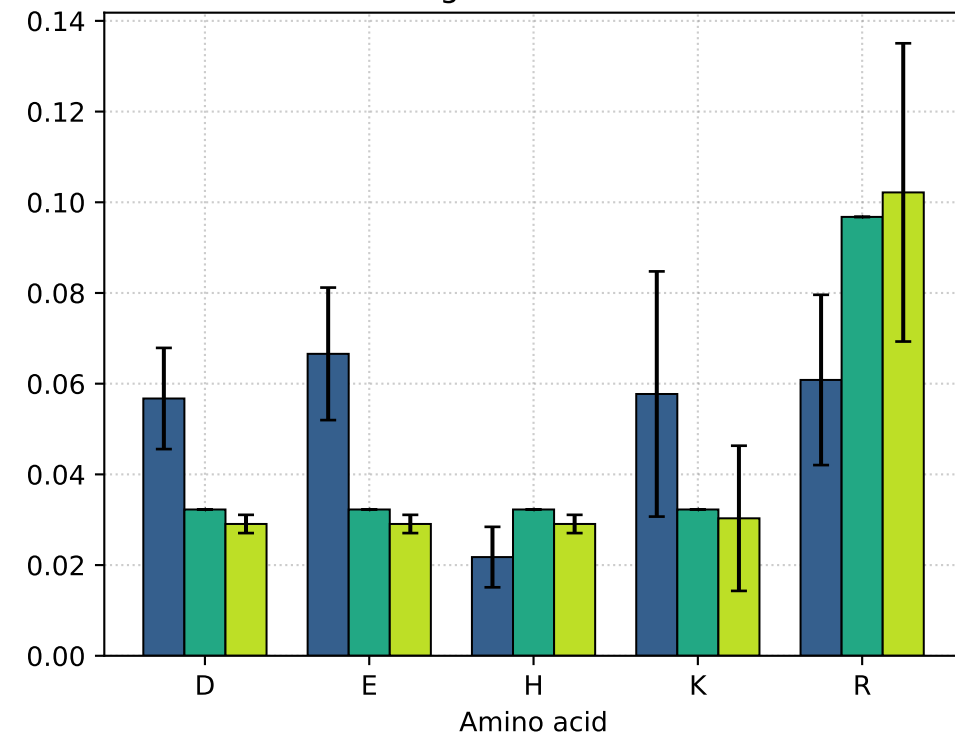
**Codon + GC correlation:**

- Mean Pearson:
- Coefficient: 0.36190
- p-value: 2.03093e-01

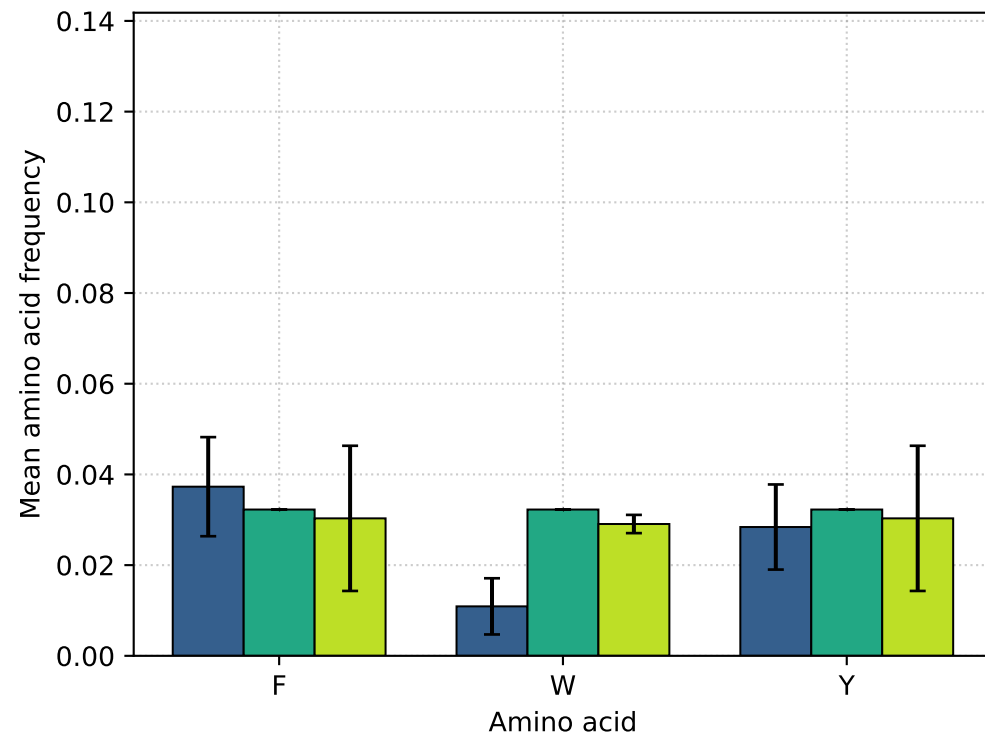
- Mean Spearman:

- Coefficient: 0.38615
- p-value: 1.36314e-01

## Charged amino acids



## Aromatic amino acids



## Uncharged amino acids

