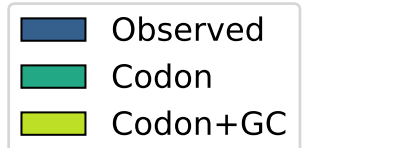
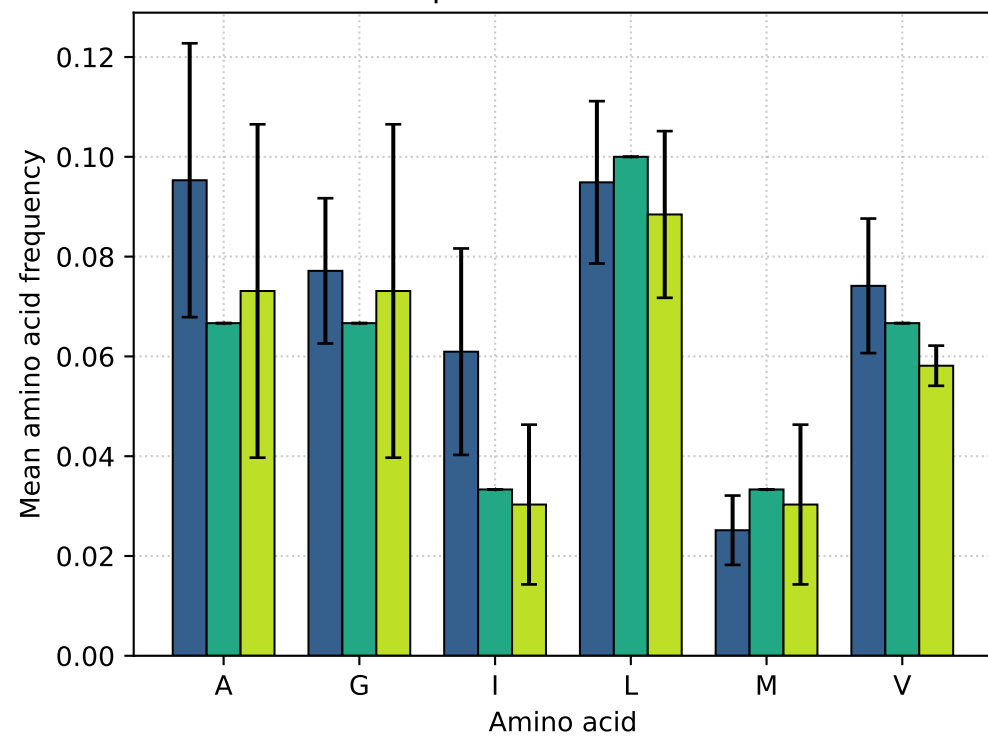


# Bacteria - Mean proteome amino acid frequency for genetic code: Vertebrate mitochondrial

## Aliphatic amino acids



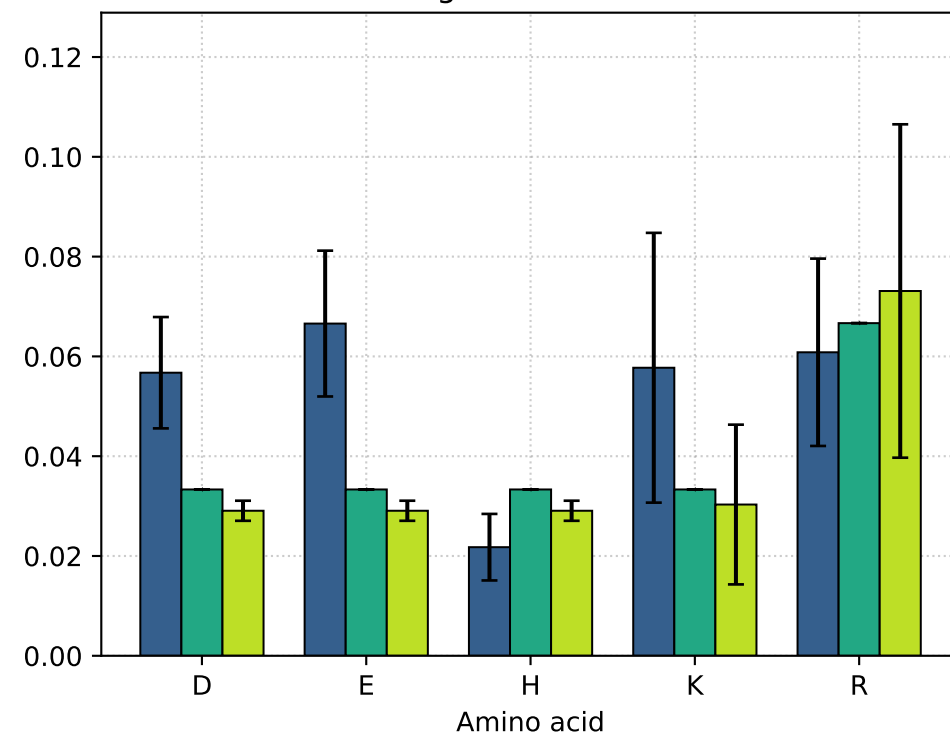
Codon correlation:

- Mean Pearson:
- Coefficient: 0.53580
- p-value: 6.15193e-02
- Mean Spearman:
- Coefficient: 0.55334
- p-value: 5.90285e-02

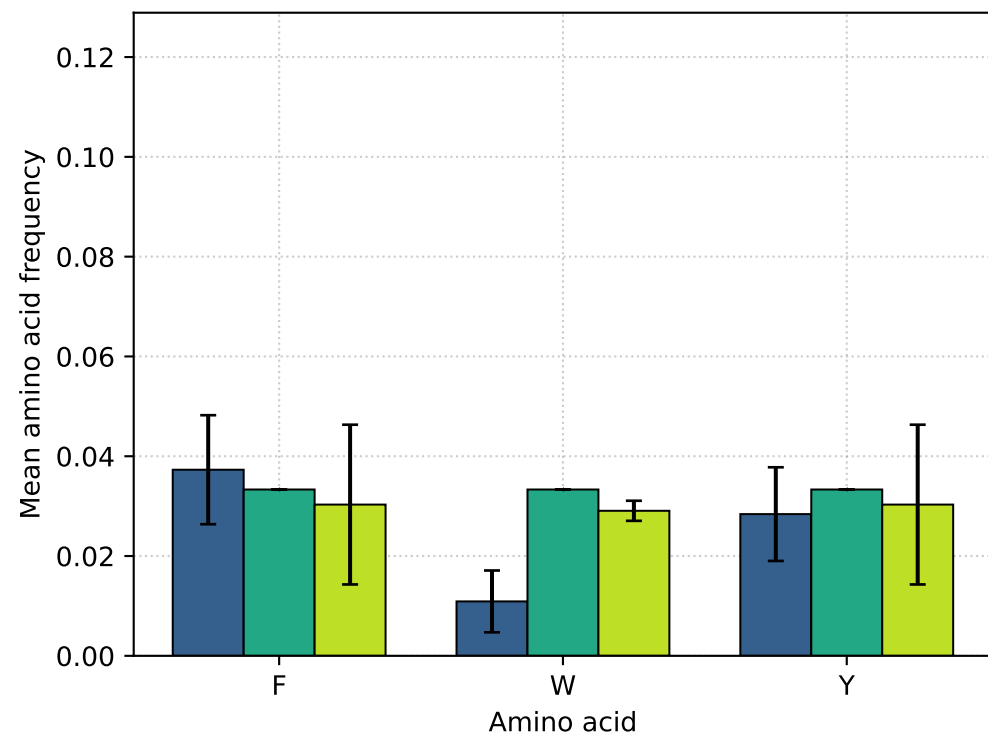
Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.58780
- p-value: 2.84809e-02
- Mean Spearman:
- Coefficient: 0.54245
- p-value: 4.24740e-02

## Charged amino acids



## Aromatic amino acids



## Uncharged amino acids

