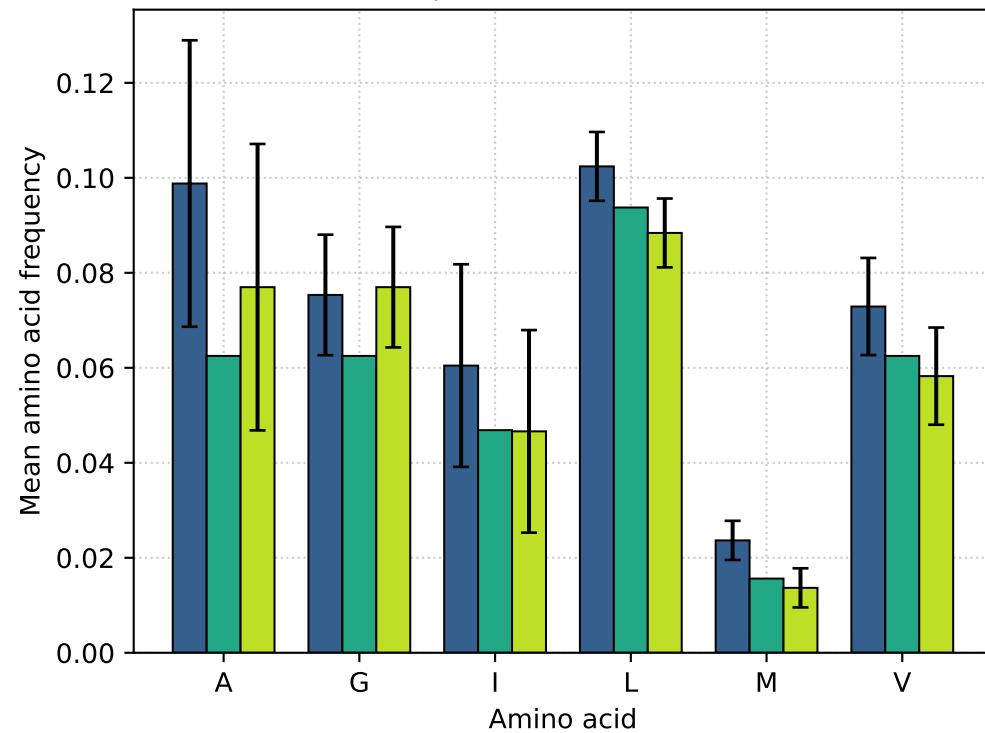


# Bacteria - Mean proteome amino acid frequency for genetic code: Blastocrithidia nuclear

## Aliphatic amino acids



Observed  
Codon  
Codon+GC

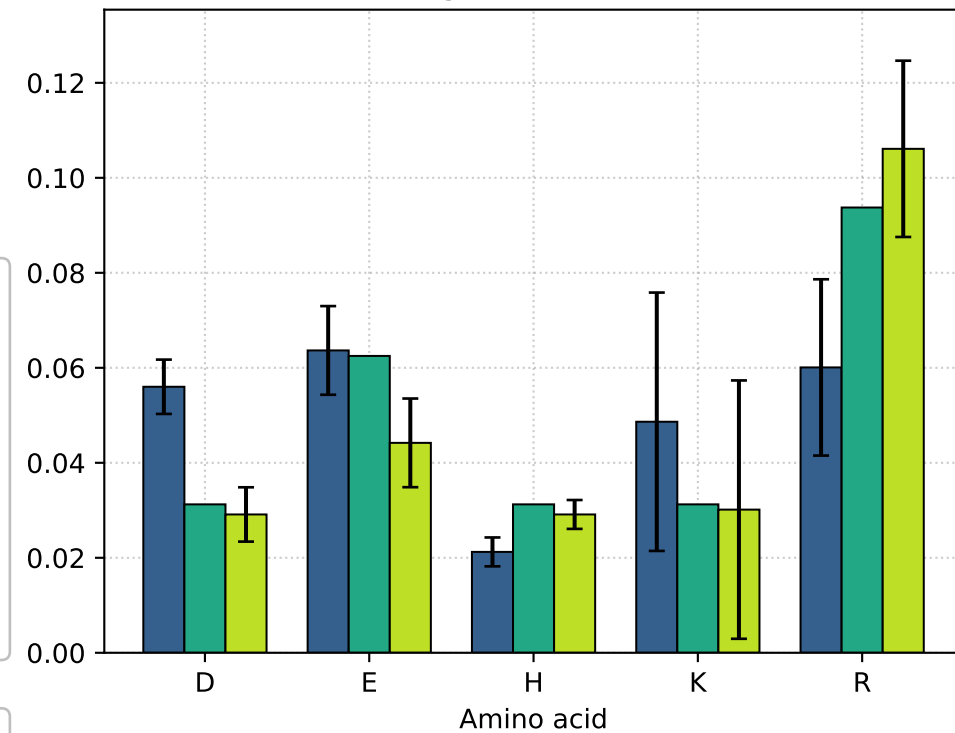
### Codon correlation:

- Mean Pearson:
- Coefficient: 0.63753
- p-value: 2.31979e-02

- Mean Spearman:
- Coefficient: 0.69198
- p-value: 1.42513e-02

- Mean log-MSE: 0.52488

## Charged amino acids



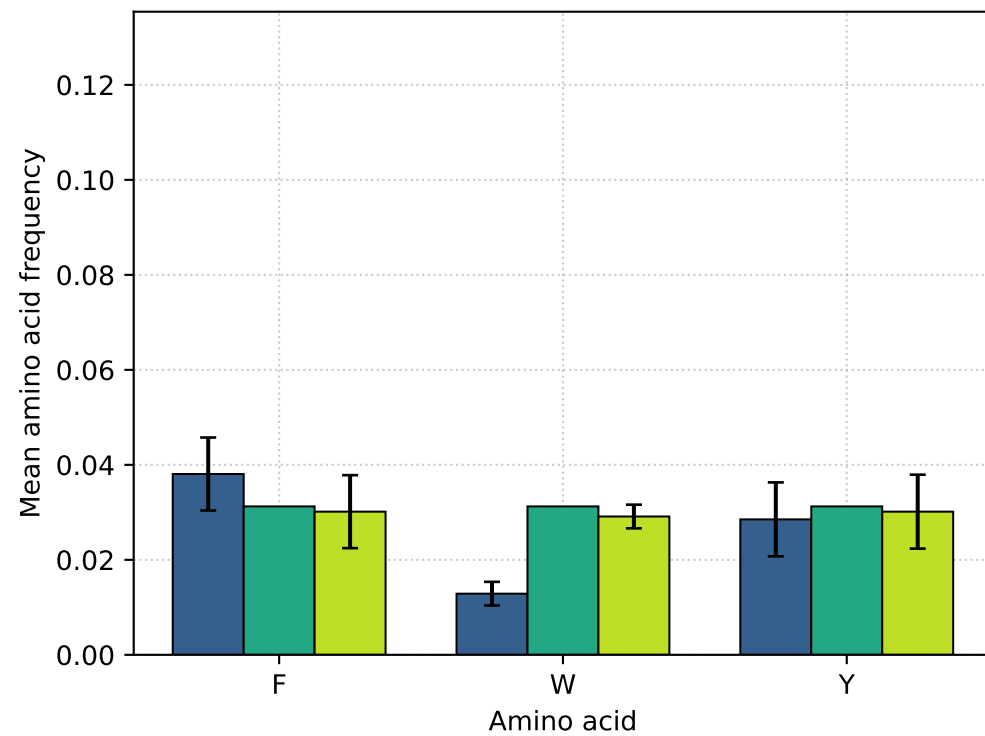
### Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.71401
- p-value: 9.02395e-04

- Mean Spearman:
- Coefficient: 0.71748
- p-value: 1.01289e-03

- Mean log-MSE: 0.58743

## Aromatic amino acids



## Uncharged amino acids

