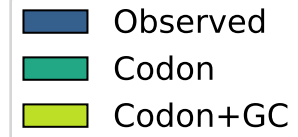
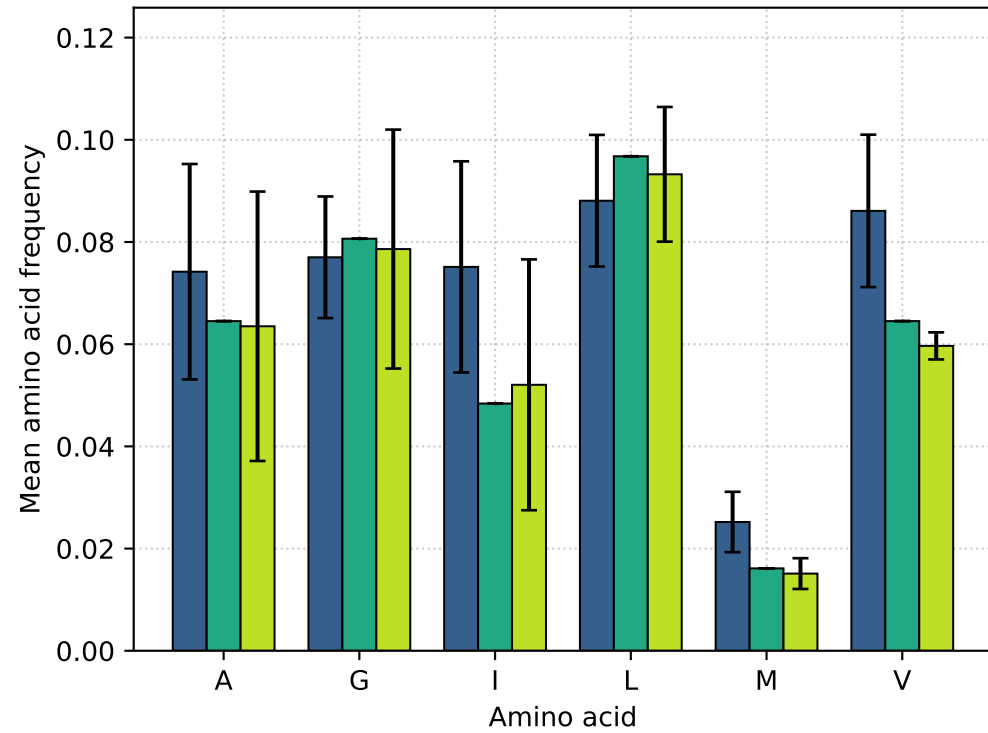


# Archaea - Mean proteome amino acid frequency for genetic code: Candidate division sr1 and gracilibacteria

## Aliphatic amino acids



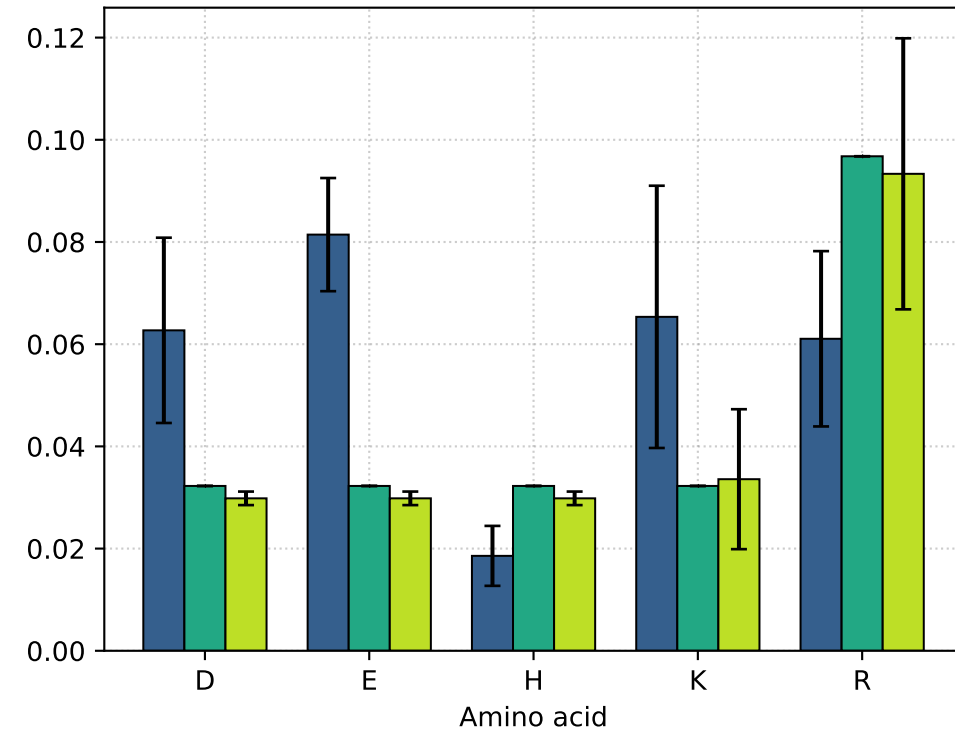
Codon correlation:  
- Mean Pearson:  
- Coefficient: 0.53881  
- p-value: 3.82092e-02

- Mean Spearman:  
- Coefficient: 0.59671  
- p-value: 1.60882e-02

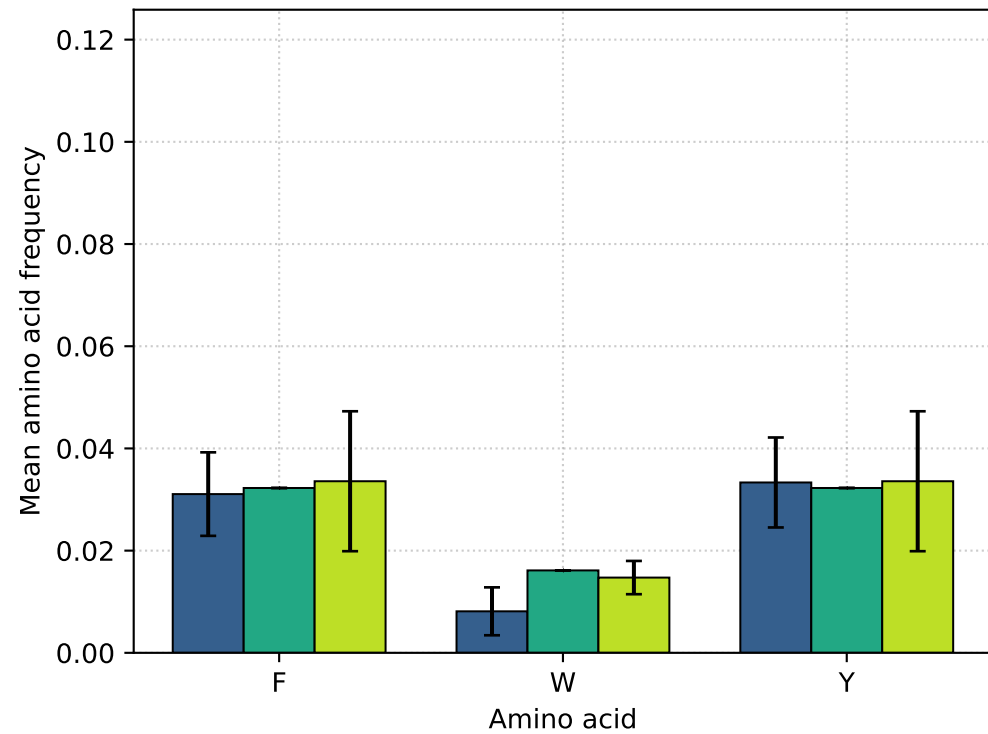
Codon + GC correlation:  
- Mean Pearson:  
- Coefficient: 0.58549  
- p-value: 2.41973e-02

- Mean Spearman:  
- Coefficient: 0.60469  
- p-value: 1.24153e-02

## Charged amino acids



## Aromatic amino acids



## Uncharged amino acids

