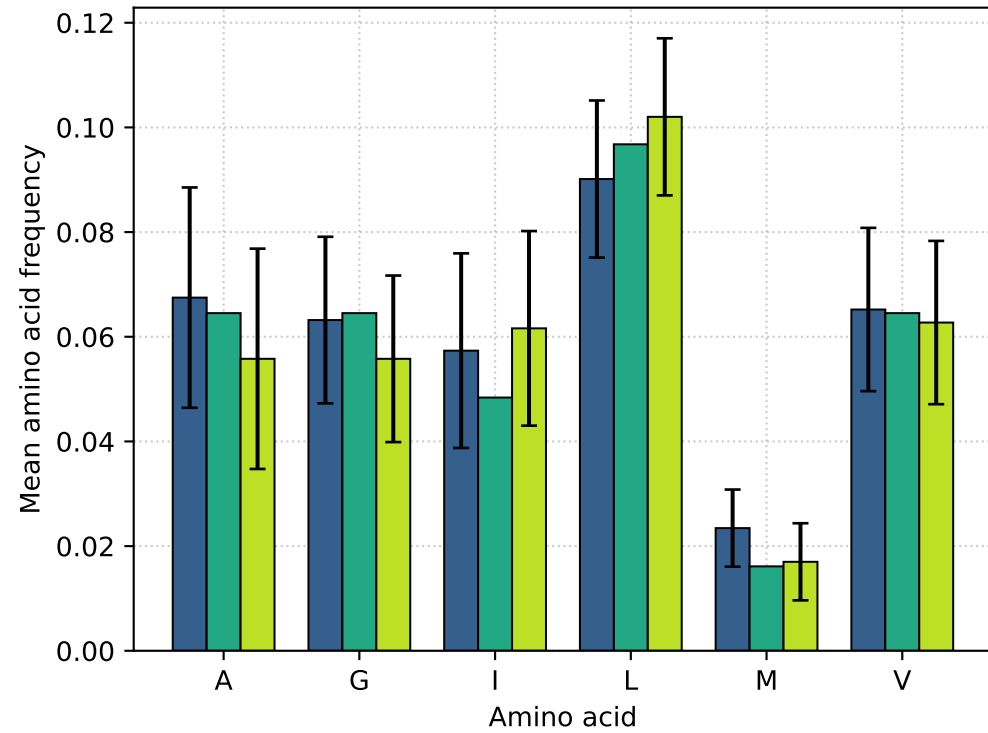


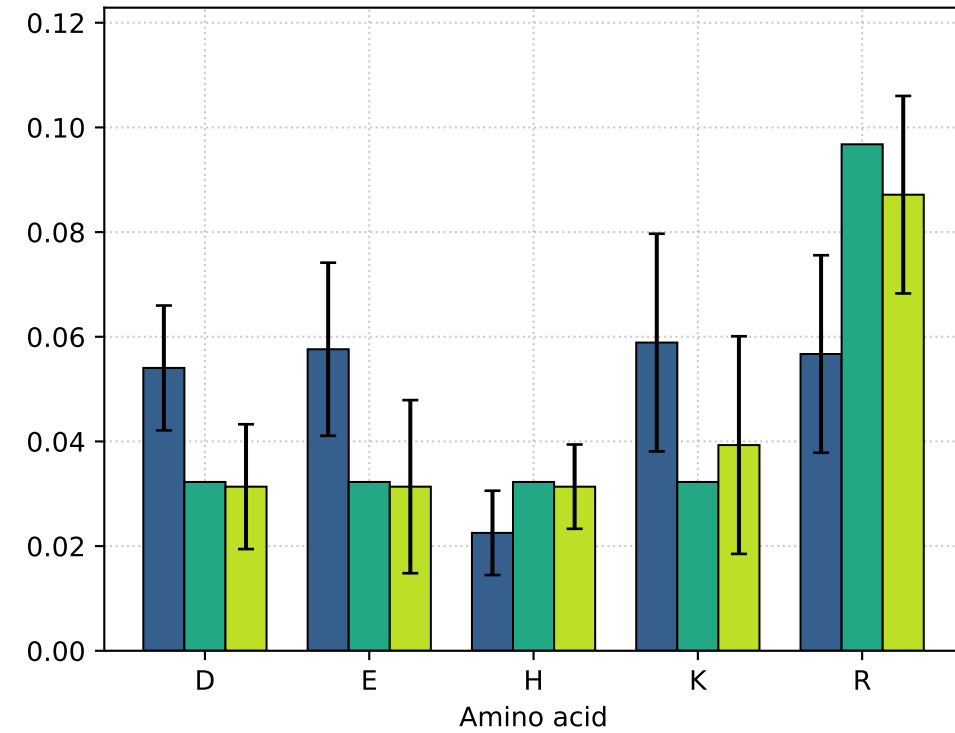
# Viruses - Mean proteome amino acid frequency for genetic code: Mold-protozoan-coelenterate mitochondrial and mycoplasma spiroplasma

## Aliphatic amino acids



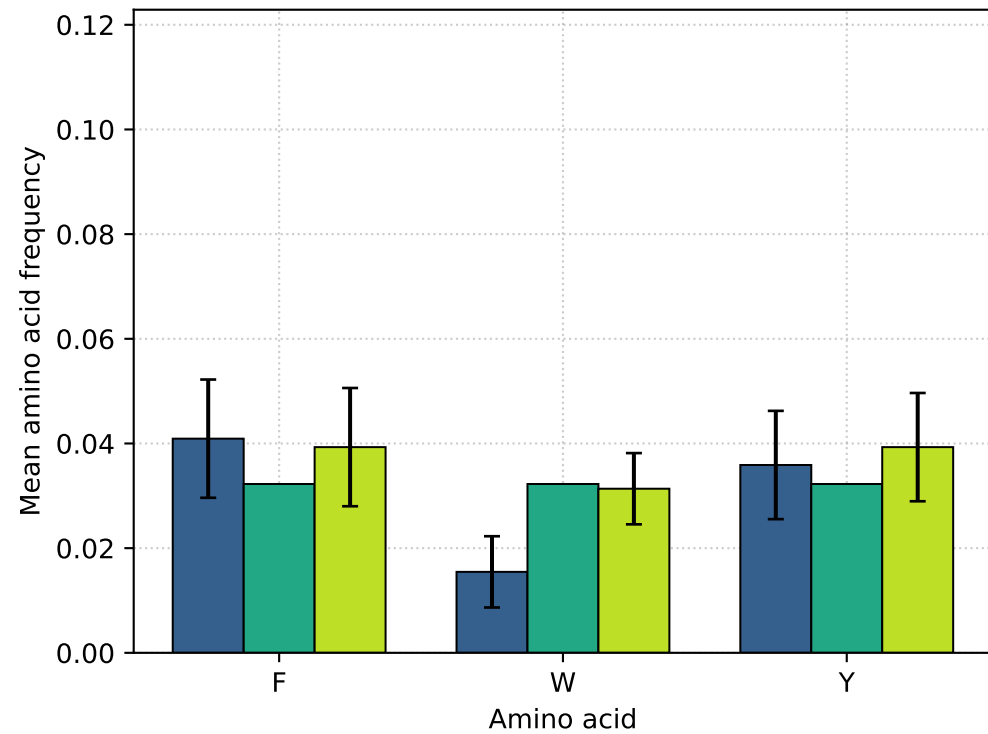
Codon correlation:  
- Mean Pearson:  
- Coefficient: 0.61050  
- p-value: 4.37991e-02  
  
- Mean Spearman:  
- Coefficient: 0.62252  
- p-value: 3.76680e-02

## Charged amino acids



Codon + GC correlation:  
- Mean Pearson:  
- Coefficient: 0.68237  
- p-value: 1.08792e-02  
  
- Mean Spearman:  
- Coefficient: 0.66165  
- p-value: 1.12080e-02

## Aromatic amino acids



## Uncharged amino acids

