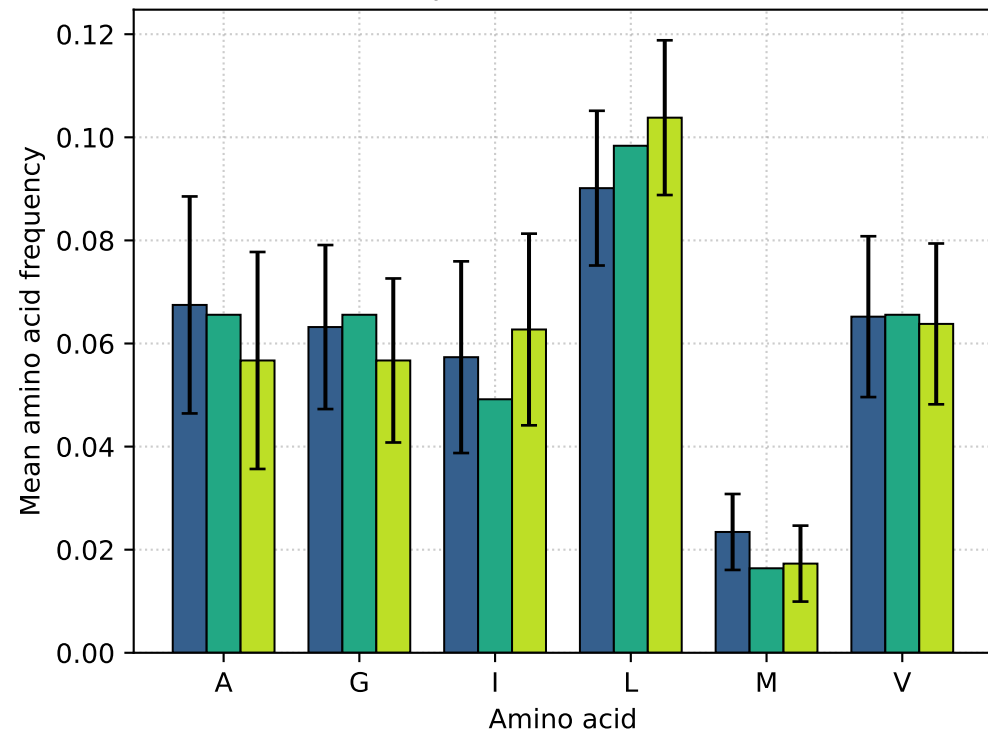


# Viruses - Mean proteome amino acid frequency for genetic code: Standard

## Aliphatic amino acids



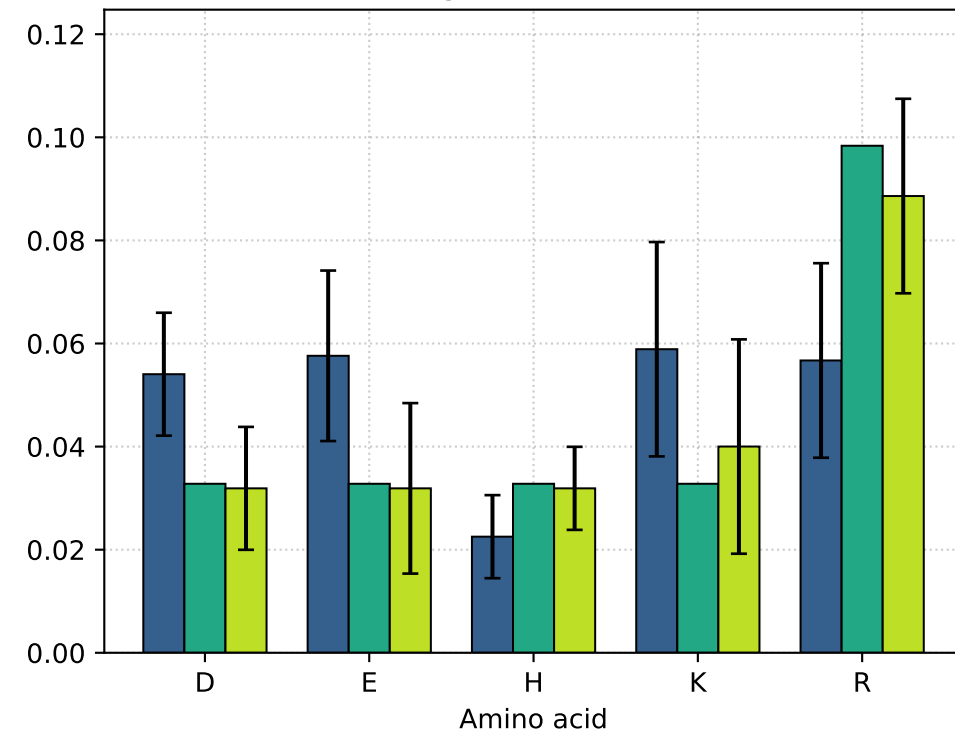
Observed  
Codon  
Codon+GC

### Codon correlation:

- Mean Pearson:  
- Coefficient: 0.63780  
- p-value: 3.40820e-02

- Mean Spearman:  
- Coefficient: 0.65762  
- p-value: 2.43372e-02

## Charged amino acids

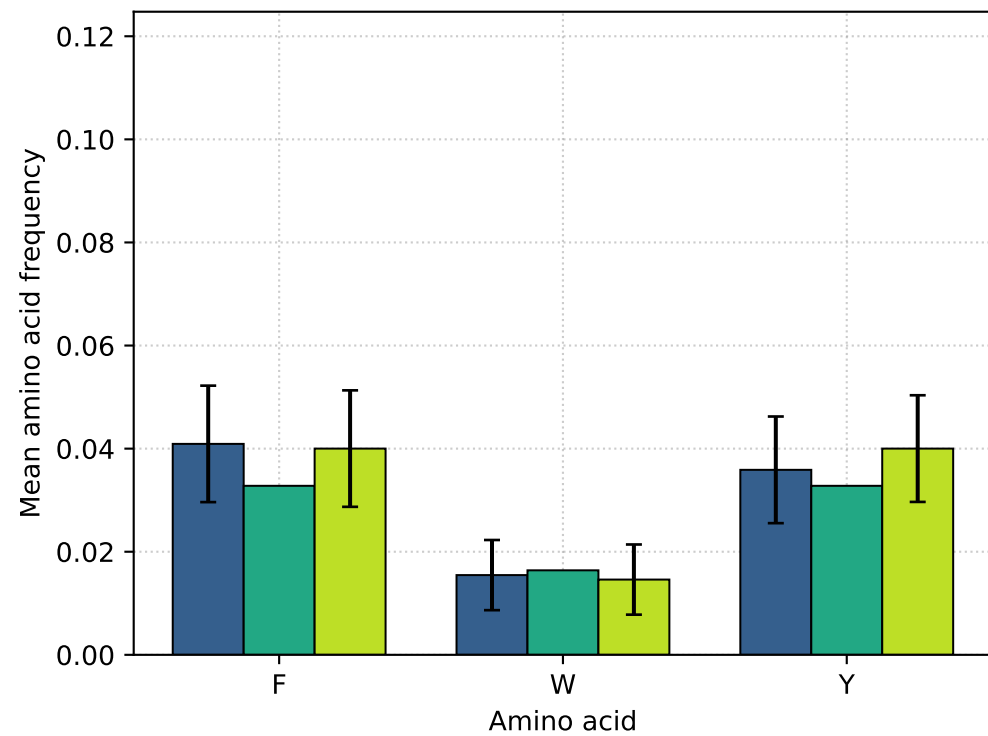


### Codon + GC correlation:

- Mean Pearson:  
- Coefficient: 0.70821  
- p-value: 7.92592e-03

- Mean Spearman:  
- Coefficient: 0.68963  
- p-value: 7.82837e-03

## Aromatic amino acids



## Uncharged amino acids

