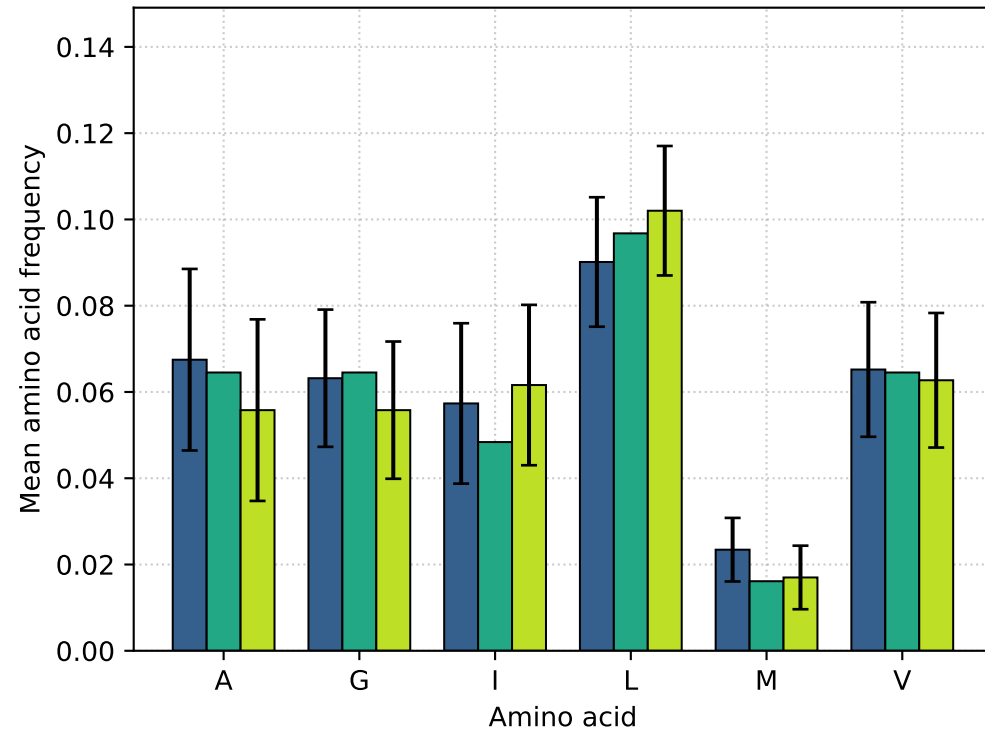


# Viruses - Mean proteome amino acid frequency for genetic code: Echinoderm-flatworm mitochondrial

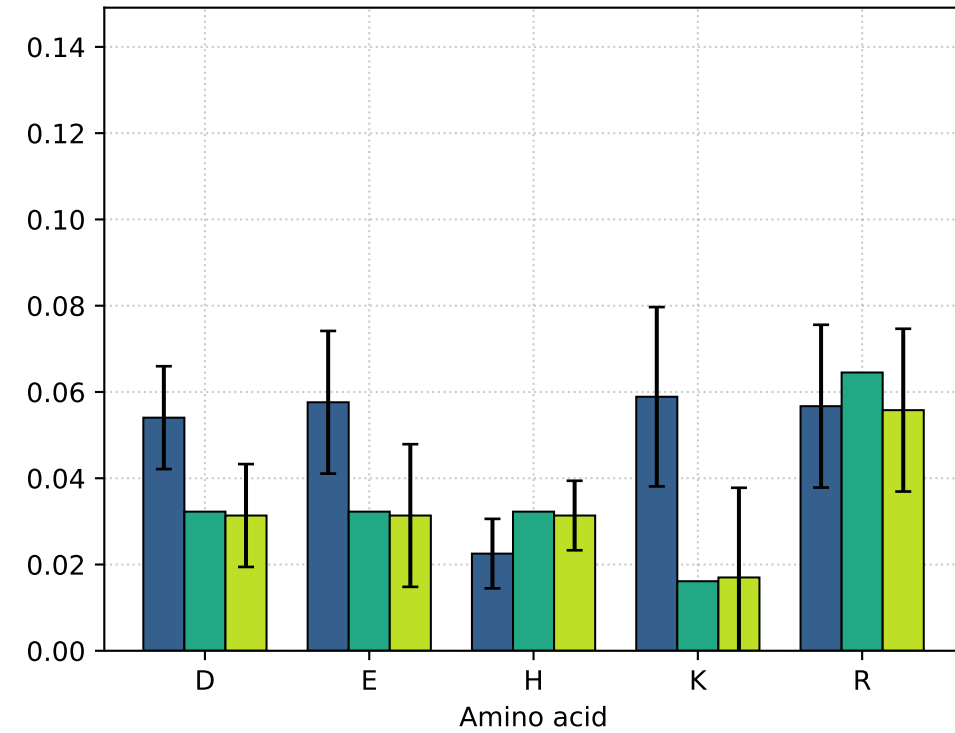
## Aliphatic amino acids



Observed  
Codon  
Codon+GC

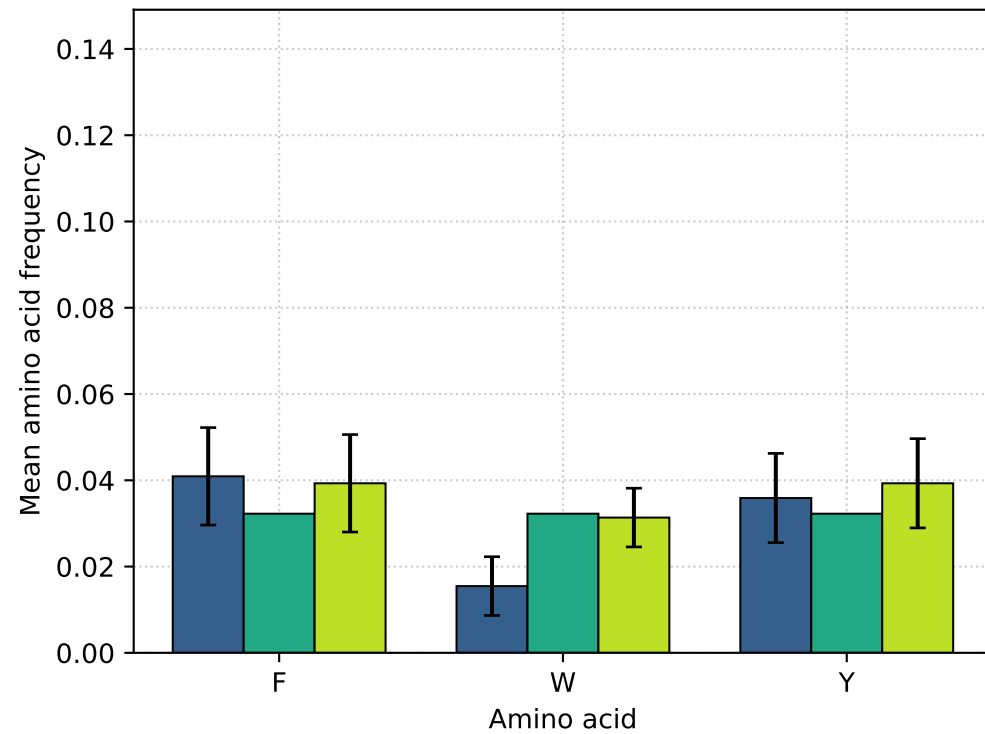
*Codon correlation:*  
- Mean Pearson:  
- Coefficient: 0.58053  
- p-value: 5.15131e-02  
  
- Mean Spearman:  
- Coefficient: 0.58471  
- p-value: 5.01417e-02

## Charged amino acids



*Codon + GC correlation:*  
- Mean Pearson:  
- Coefficient: 0.61646  
- p-value: 2.95517e-02  
  
- Mean Spearman:  
- Coefficient: 0.58511  
- p-value: 3.63786e-02

## Aromatic amino acids



## Uncharged amino acids

