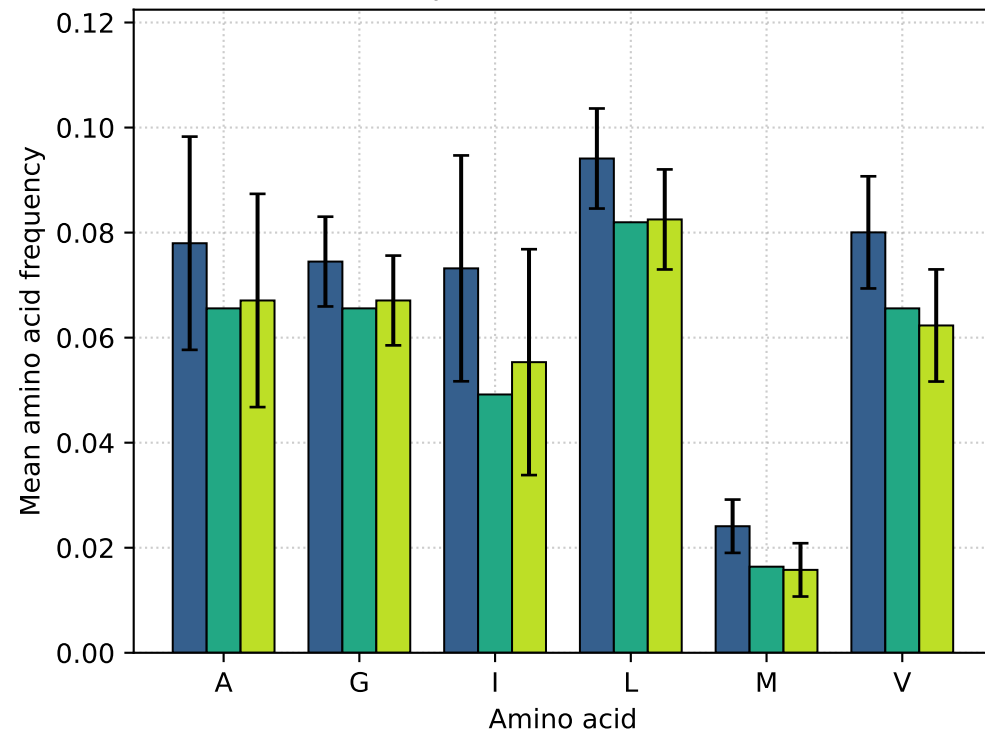


# Archaea - Mean proteome amino acid frequency for genetic code: Alternative yeast

## Aliphatic amino acids



### Codon correlation:

- Mean Pearson:  
- Coefficient: 0.50674  
- p-value: 5.61321e-02

- Mean Spearman:  
- Coefficient: 0.60377  
- p-value: 1.75392e-02

- Mean log-MSE: 0.61800

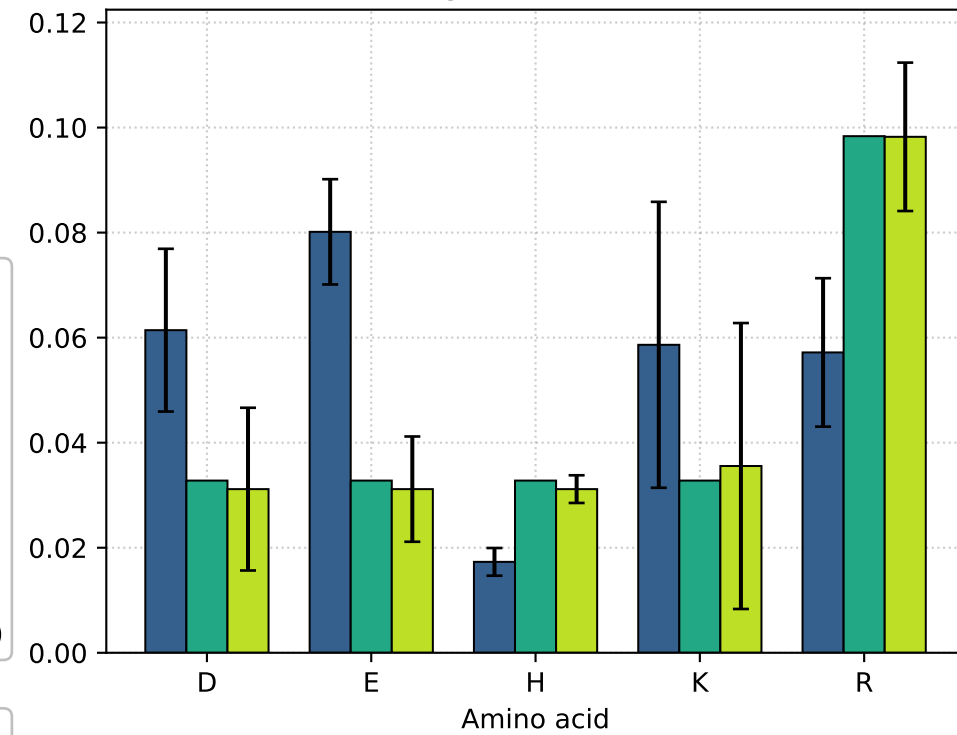
### Codon + GC correlation:

- Mean Pearson:  
- Coefficient: 0.59500  
- p-value: 9.25052e-03

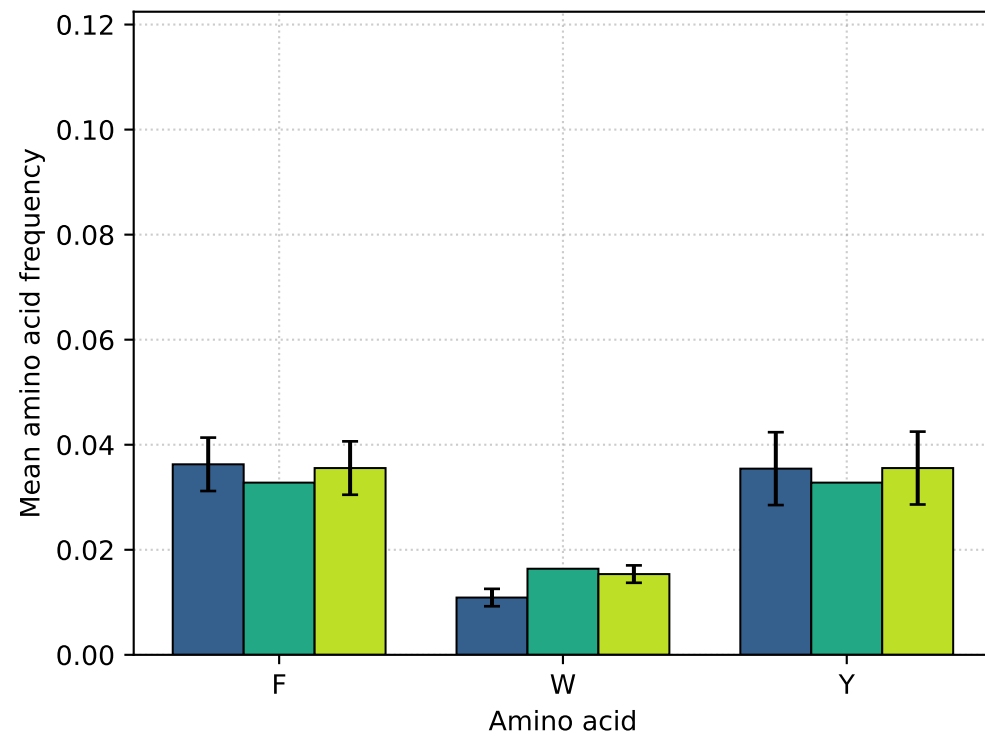
- Mean Spearman:  
- Coefficient: 0.62889  
- p-value: 4.59777e-03

- Mean log-MSE: 0.62009

## Charged amino acids



## Aromatic amino acids



## Uncharged amino acids

