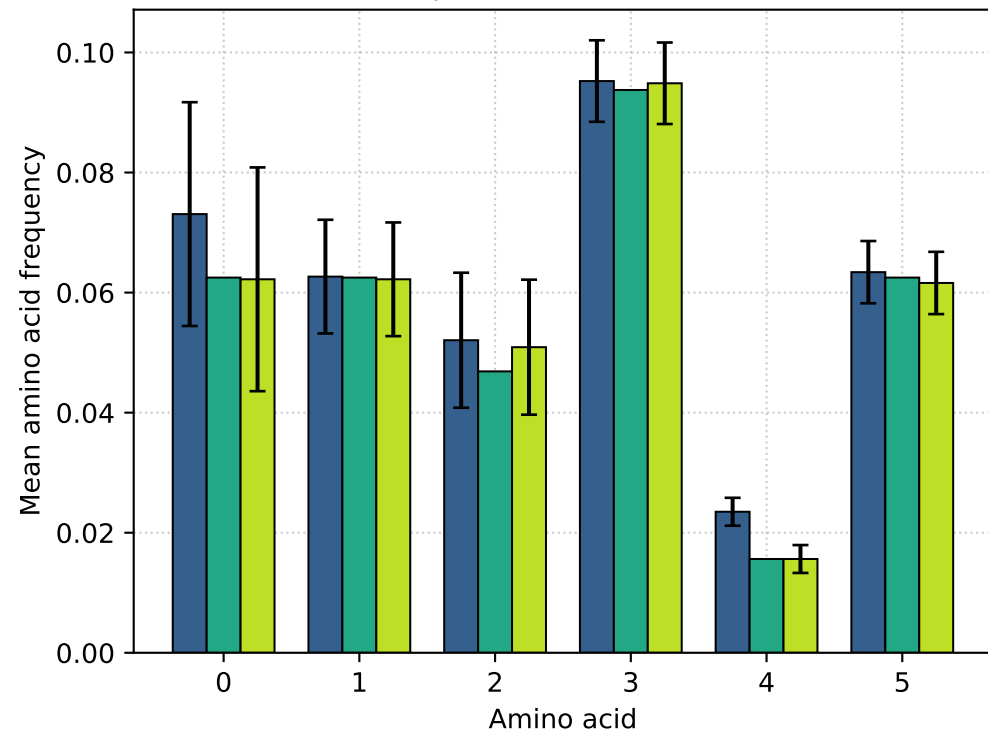
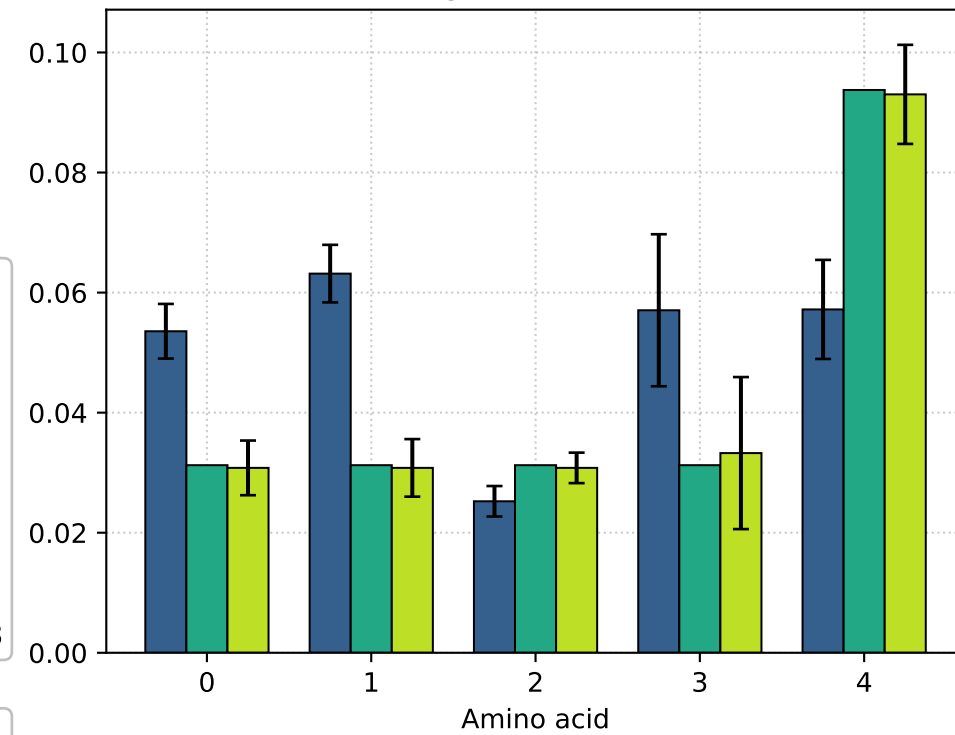


# Eukaryota - Mean proteome amino acid frequency for genetic code: Karyorelict nuclear

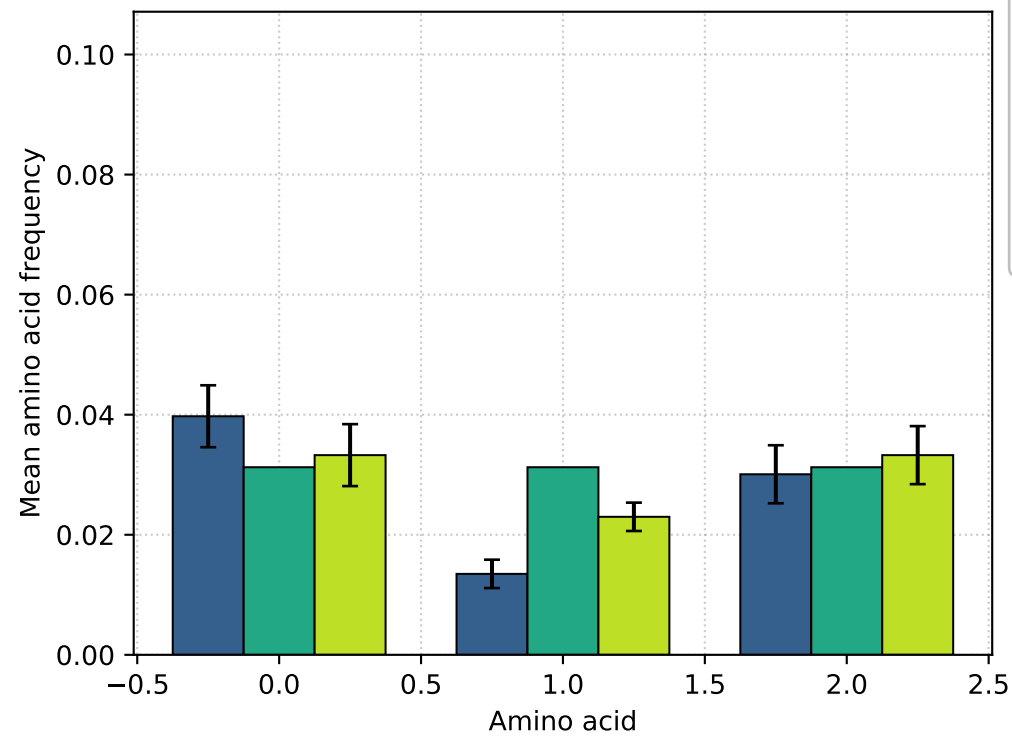
## Aliphatic amino acids



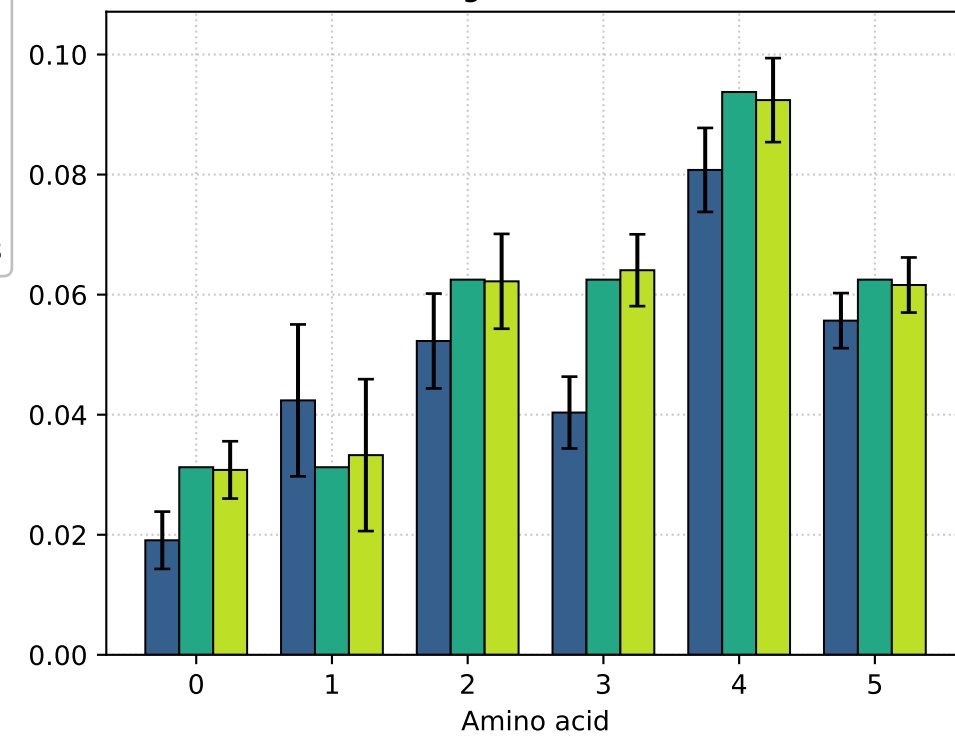
## Charged amino acids



## Aromatic amino acids



## Uncharged amino acids



### Codon correlation:

- Mean Pearson:
- Coefficient: 0.69782
- p-value: 1.68519e-02

- Mean Spearman:
- Coefficient: 0.67630
- p-value: 2.28897e-02

- Mean log-MSE: 0.36868

### Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.75041
- p-value: 5.15894e-04

- Mean Spearman:
- Coefficient: 0.72230
- p-value: 1.79354e-03

- Mean log-MSE: 0.31423