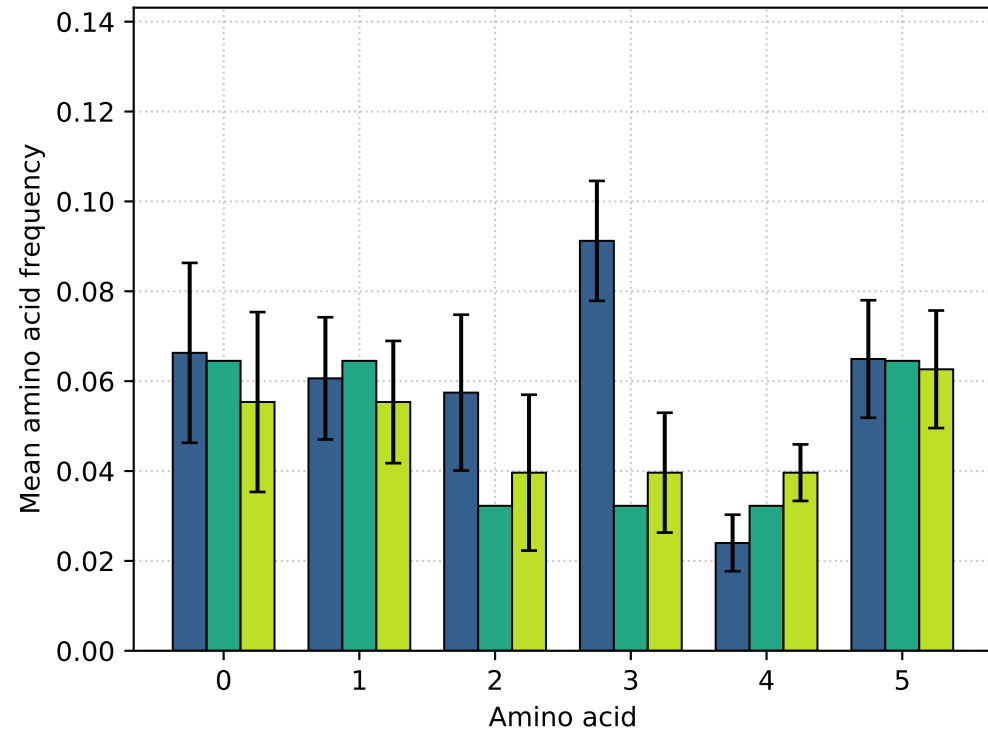
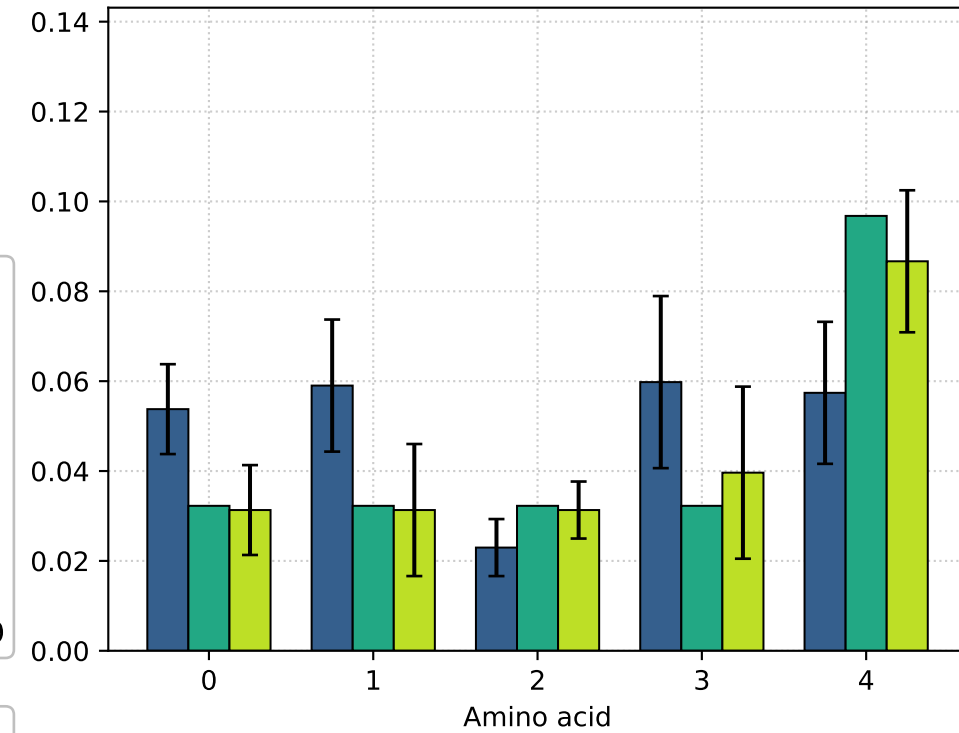


# Viruses - Mean proteome amino acid frequency for genetic code: Yeast mitochondrial

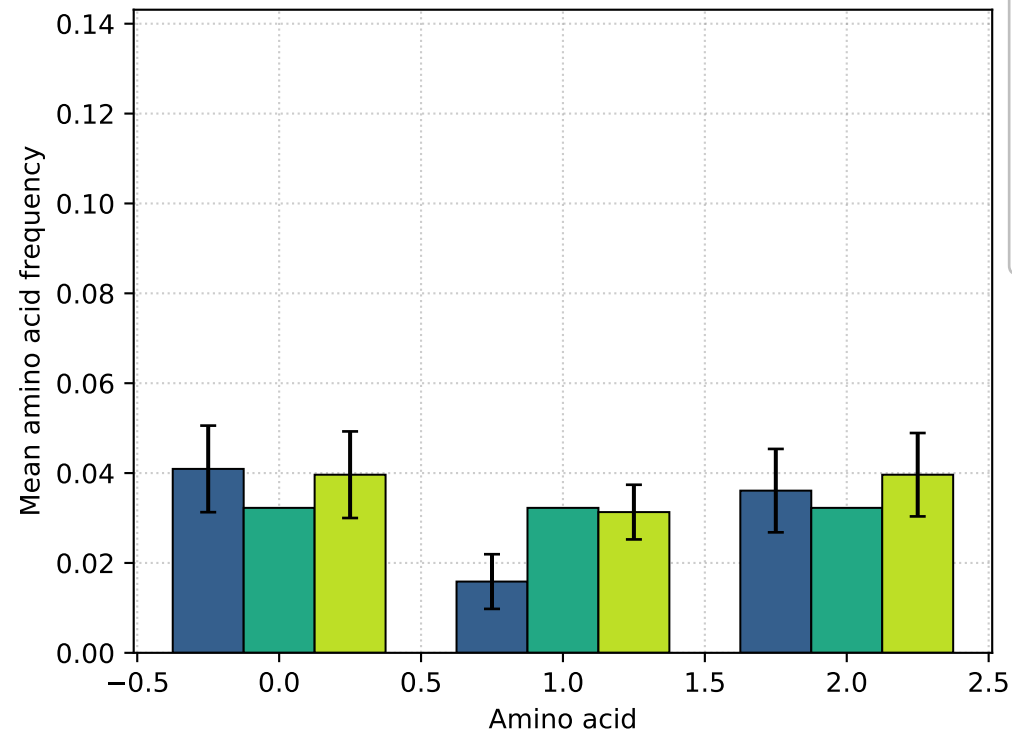
## Aliphatic amino acids



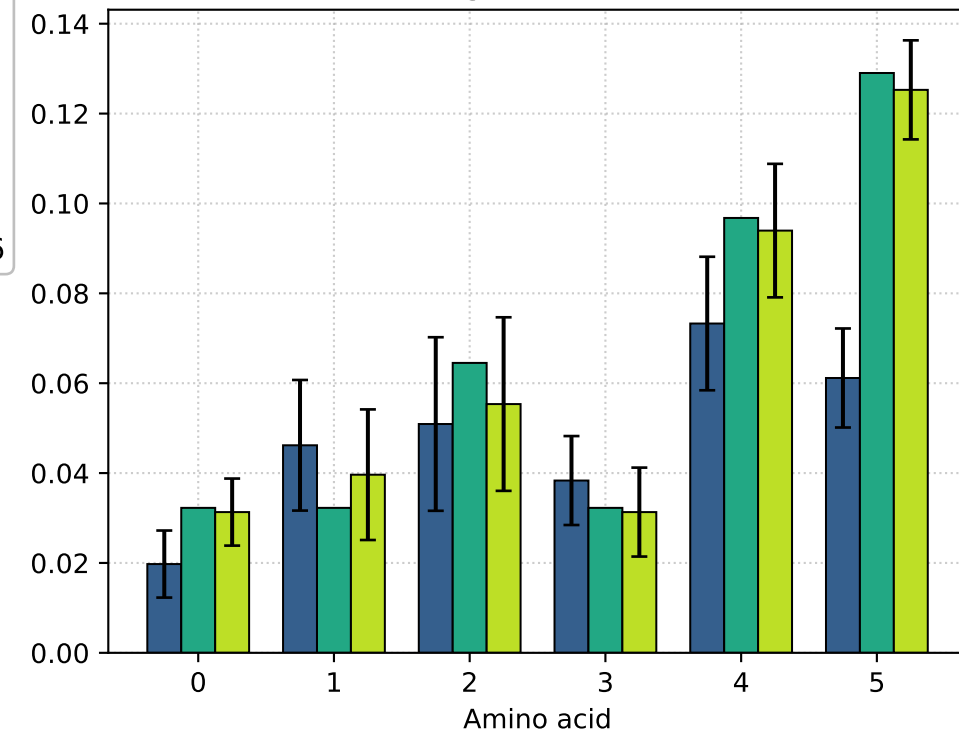
## Charged amino acids



## Aromatic amino acids



## Uncharged amino acids



**Codon correlation:**

- Mean Pearson:
- Coefficient: 0.34998
- p-value: 2.12021e-01
- Mean Spearman:
- Coefficient: 0.41623
- p-value: 1.77649e-01
- Mean log-MSE: 0.64699

**Codon + GC correlation:**

- Mean Pearson:
- Coefficient: 0.43229
- p-value: 1.02713e-01
- Mean Spearman:
- Coefficient: 0.49258
- p-value: 5.95849e-02
- Mean log-MSE: 0.56446