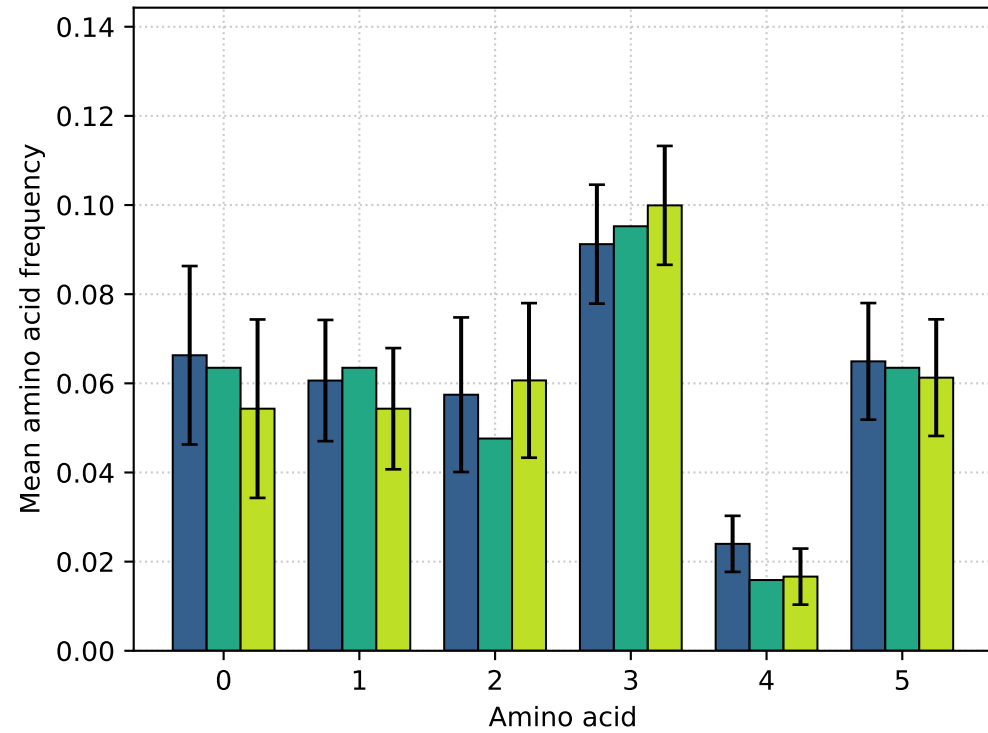


# Viruses - Mean proteome amino acid frequency for genetic code: Alternative flatworm mitochondrial

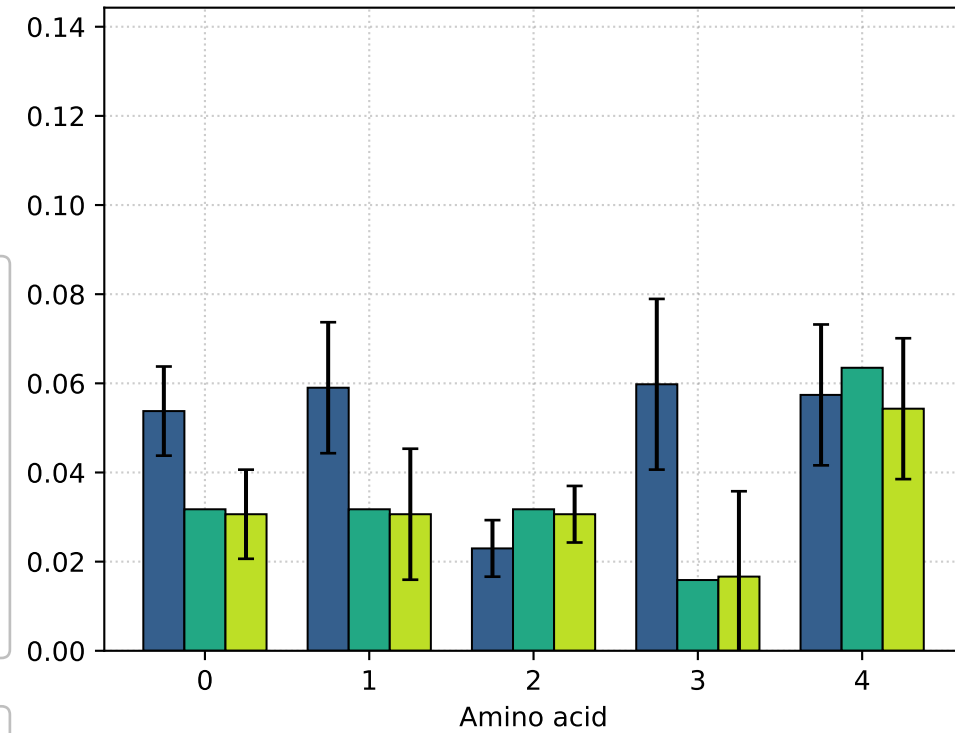
## Aliphatic amino acids



Codon correlation:

- Mean Pearson:
- Coefficient: 0.57486
- p-value: 4.83336e-02
- Mean Spearman:
- Coefficient: 0.57039
- p-value: 5.20517e-02
- Mean log-MSE: 0.57843

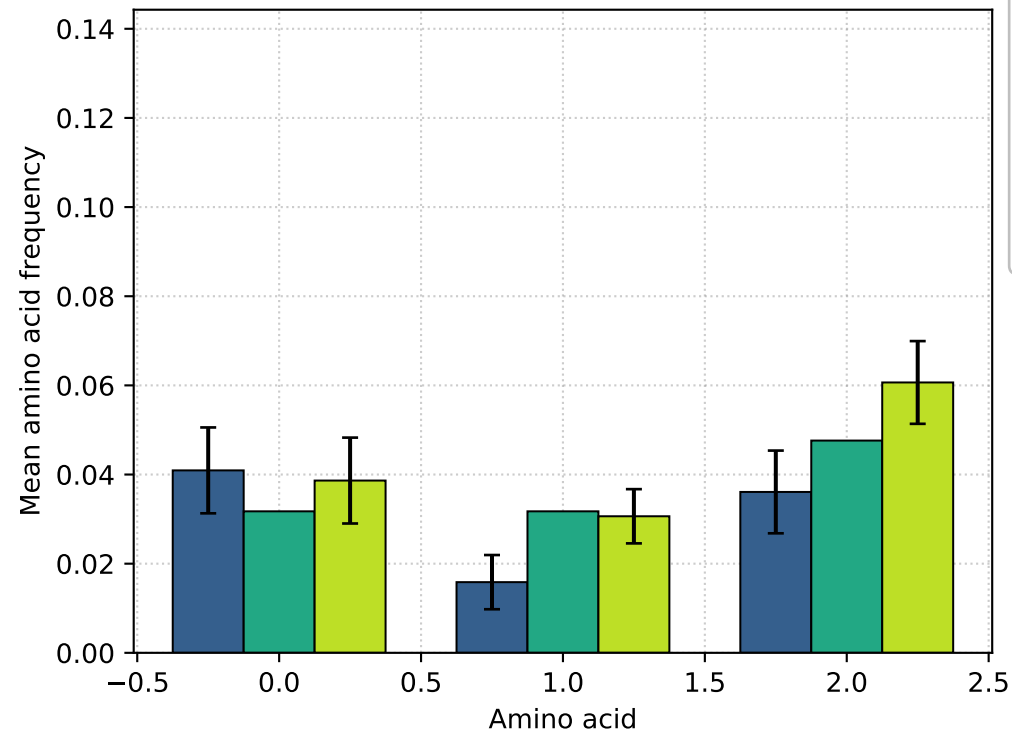
## Charged amino acids



Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.60826
- p-value: 2.67356e-02
- Mean Spearman:
- Coefficient: 0.56398
- p-value: 4.12055e-02
- Mean log-MSE: 0.56747

## Aromatic amino acids



## Uncharged amino acids

