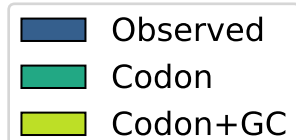
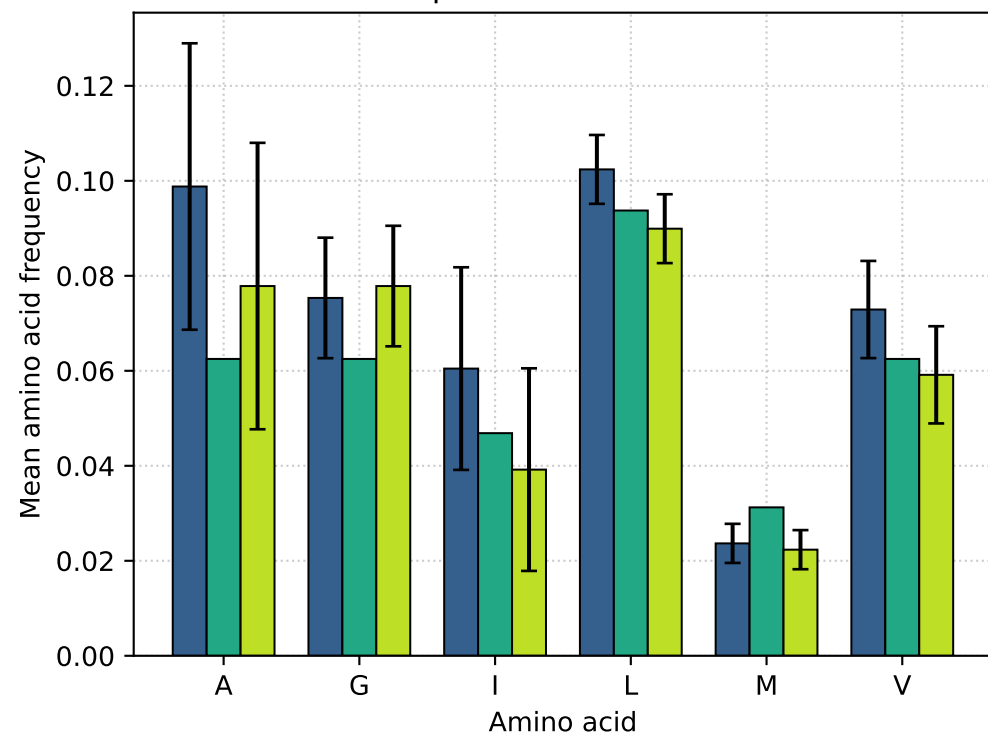


# Bacteria - Mean proteome amino acid frequency for genetic code: Rhabdopleuridae mitochondrial

## Aliphatic amino acids



### Codon correlation:

- Mean Pearson:
- Coefficient: 0.58498
- p-value: 2.20159e-02

- Mean Spearman:
- Coefficient: 0.64138
- p-value: 1.54905e-02

- Mean log-MSE: 0.56484

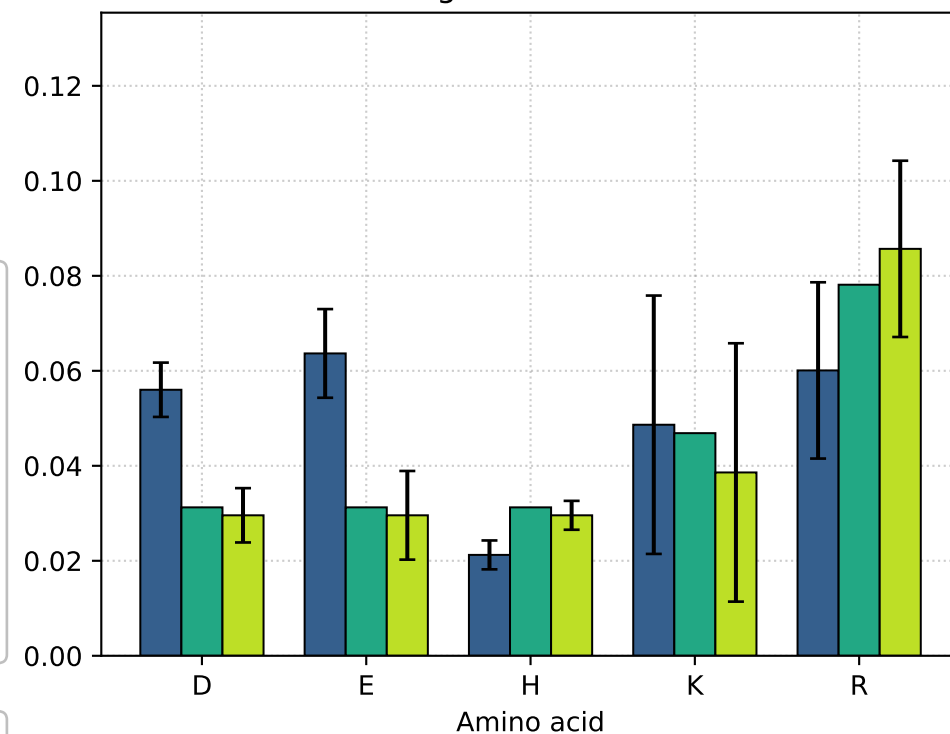
### Codon + GC correlation:

- Mean Pearson:
- Coefficient: 0.67961
- p-value: 2.68519e-03

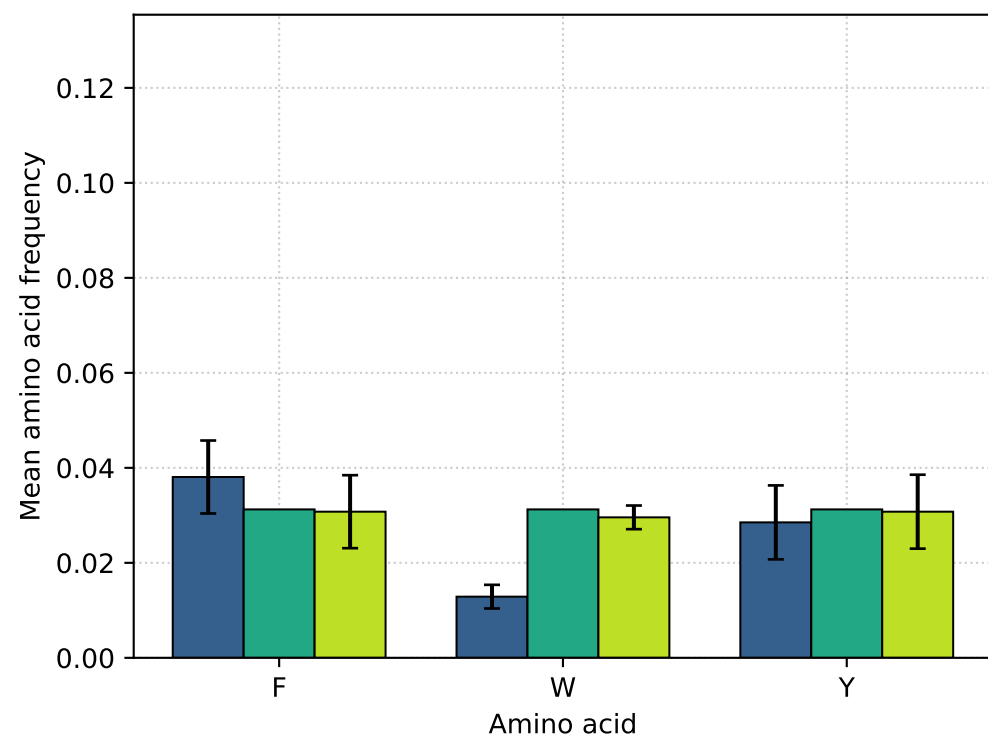
- Mean Spearman:
- Coefficient: 0.66273
- p-value: 3.86353e-03

- Mean log-MSE: 0.60528

## Charged amino acids



## Aromatic amino acids



## Uncharged amino acids

