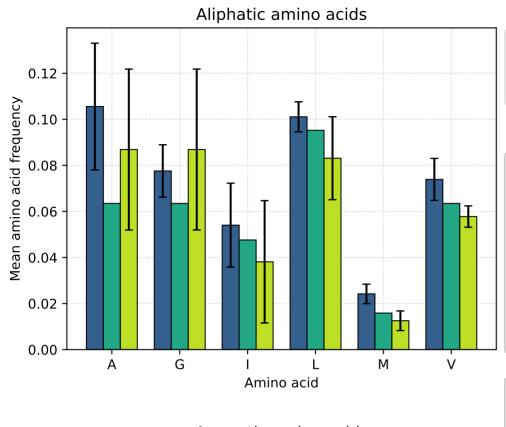
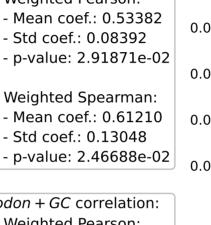
Bacteria - Weighted mean proteome amino acid frequency for genetic code: Alternative flatworm mitochondrial

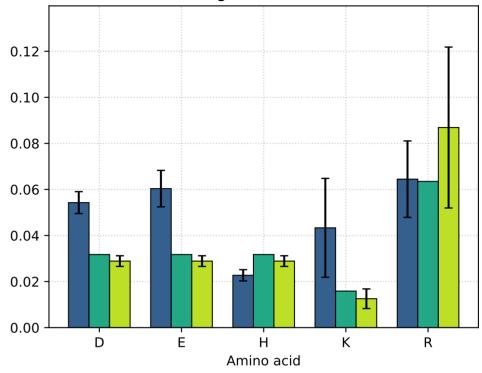




Codon correlation:

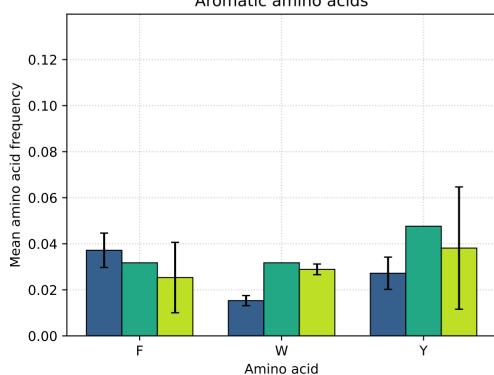
- Weighted Pearson:
- Mean coef.: 0.53382
- Std coef.: 0.08392
- Weighted Spearman:
- Mean coef.: 0.61210
- Std coef.: 0.13048
- p-value: 2.46688e-02





Charged amino acids





Codon + GC correlation:

- Weighted Pearson:
- Mean coef.: 0.64214
- Std coef.: 0.11361
- p-value: 1.09260e-02
- Weighted Spearman:
- Mean coef.: 0.62198 - Std coef.: 0.13664
- p-value: 2.15653e-02

