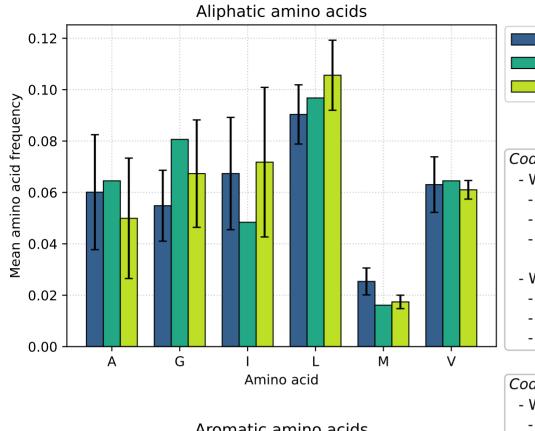
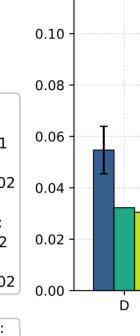
Viruses - Weighted mean proteome amino acid frequency for genetic code: Candidate division sr1 and gracilibacteria



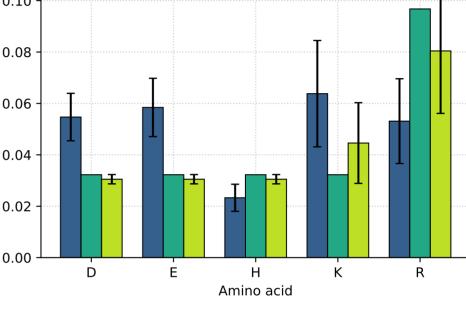


Codon correlation:

- Weighted Pearson:
- Mean coef.: 0.57141 - Std coef.: 0.22471
- p-value: 7.70201e-02
- Weighted Spearman:
- Mean coef.: 0.59452
- Std coef.: 0.21043
- p-value: 5.24817e-02



0.12



Charged amino acids

Codon + GC correlation:

- Weighted Pearson:
- Mean coef.: 0.76912
- Std coef.: 0.09413
- p-value: 1.61489e-03
- Weighted Spearman:
 - Mean coef.: 0.74165
 - Std coef.: 0.09754
 - p-value: 2.27331e-03

