Viruses - Pairwise comparison between amino acid and factors for amino acid **A** and genetic code: Candidate division sr1 and gracilibacteria 3.5 log10-Length 5.7 0.8 2.0 0.6 GC 0.4 0.2 0.125 0.100 · 0.075 · 0.050 · 0.050 · 0.050 0.025 0.125 0.100 0.075 0.050 0.025 0.10 0.15 0.6 0.025 0.050 0.075 0.100 0.125 0.05 2.0 3.5 0.2 0.4 0.025 0.050 0.075 0.100 0.125 2.5 3.0 Amino acid frequency log10-Length Codon+GC Codon+GC+Cost GC