Bacteria - Pairwise comparison between amino acid and factors for amino acid ${\bf K}$ and genetic code: Candidate division sr1 and gracilibacteria 3.0 log10-Length 0.7 5.7 0.6 GC 0.4 0.2 0.08 0.06 0.04 0.04 0.02 0.08 0.06 0.04 0.04 0.02 0.10 0.2 0.02 0.06 0.00 0.05 0.15 2.0 2.5 0.4 0.6 0.04 0.06 0.08 0.02 0.04 3.0 0.08 log10-Length GC Codon+GC Codon+GC+Cost Amino acid frequency