Archaea - Pairwise comparison between amino acid and factors for amino acid K and genetic code: Candidate division sr1 and gracilibacteria 2.8 log10-Length 7.7 9.7 9.7 9.7 2.2 0.6 设 0.5 0.4 0.3 0.06 0.05 0.03 29 + 0.04 0.03 0.02 -0.06 0.06 0.05 0.04 0.03 0.02 0.02 0.03 0.04 0.05 0.02 0.04 0.06 0.08 0.10 0.122.2 2.6 2.8 0.3 0.02 0.03 0.04 0.05 0.06 0.4 0.5 0.6 GC Codon+GC Codon+GC+Cost Amino acid frequency log10-Length