Archaea - Pairwise comparison between amino acid and factors for amino acid **H** and genetic code: Condylostoma nuclear 2.8 log10-Length 7.7 9.7 2.2 -0.6 设 0.5 0.4 0.3 0.031 -0.030 0.029 0.028 0.027 0.032 to 0.031 0.030 0.029 0.029 0.028 0.02 0.03 0.04 2.2 2.6 2.8 0.3 0.6 0.027 0.028 0.029 0.030 0.031 0.028 0.029 0.030 0.031 0.032 0.01 2.4 0.4 0.5 Amino acid frequency log10-Length GC Codon+GC Codon+GC+Cost