Exogenous and Combined Codon Usage -1.0 -0.5 0.0 0.5 1.0 -1.0 -0.5 0.0 0.5-1.0 -0.5 0.0 -1.0 -0.5 0.0GCA GCC GCG GCT 8.0 0.8 0.6 0.6 0.4 0.4 0.2 0.2 0.0 0.0 GGA GGC GGG GGT CAC CAT ATA ATC ATT TTC TTT G Н F 1.0 1.0 8.0 0.8 0.6 0.6 0.4 0.4 0.2 0.2 0.0 0.0 CTA CTC CTG CTT TTA AAC AAT CCA CCC CCG CCT Κ Ν 1.0 1.0 **Proportion** 8.0 0.8 0.6 0.6 0.4 0.4 0.2 0.2 0.0 0.0 TCA TCC TCG TCT CAA CAG AGA AGG CGA CGC CGG R S ACA 1.0 1.0 ACC ACG ACT 8.0 0.8 0.6 0.6 0.4 0.4 0.2 0.2 0.0 0.35 GTA GTC GTG GTT TAC TAT AGC AGT Z 1.0 Model fit - · Exogenous · · · Combined 0.30 8.0 0.25 0.6 0.15 0.4 0.10 0.2 0.05 0.0 0.5 1.0 -1.0 -0.5 0.0 -1.0 -0.5 -1.0 -0.50.0 1.0 0.0 0.5 log₁₀ (Protein Synthesis Rate ♦)