k_Archaea;p_Miscellaneous Crenarchaeotic Group I k__Archaea;p__Thaumarchaeota k_Archaea;p_Woesearchaeota (DHVEG-6) k Bacteria;Other k_Bacteria;p_Acidobacteria k Bacteria;p Actinobacteria k Bacteria;p Aerophobetes k Bacteria;p Armatimonadetes k_Bacteria;p_Bacteroidetes ■ k_Bacteria;p_Candidate division OP3 k_Bacteria;p_Candidate division SR1 k_Bacteria;p_Candidate division WS6 k_Bacteria;p_Chlamydiae k_Bacteria;p_Chlorobi k_Bacteria;p_Chloroflexi k_Bacteria;p_Cyanobacteria k_Bacteria;p_Deferribacteres k Bacteria;p Deinococcus-Thermus k_Bacteria;p_Elusimicrobia k Bacteria;p Fibrobacteres k_Bacteria;p_Firmicutes k__Bacteria;p__Fusobacteria k Bacteria;p GAL08 k Bacteria;p Gemmatimonadetes ■ k__Bacteria;p__Gracilibacteria k_Bacteria;p_Hydrogenedentes k_Bacteria;p_JL-ETNP-Z39 k_Bacteria;p_Latescibacteria k_Bacteria;p_Lentisphaerae k Bacteria;p Microgenomates k Bacteria; p Nitrospirae k_Bacteria;p_Omnitrophica k_Bacteria;p_Parcubacteria k_Bacteria;p_Planctomycetes k Bacteria;p Proteobacteria k_Bacteria;p_SHA-109 k_Bacteria;p_SM2F11 k_Bacteria;p_Saccharibacteria k_Bacteria;p_Spirochaetae k_Bacteria;p_Synergistetes k_Bacteria;p_TM6 k_Bacteria;p_Tenericutes k_Bacteria;p_Thermotogae

k_Bacteria;p_Verrucomicrobia k_Bacteria;p_WCHB1-60 k_Bacteria;p_WD272

Unassigned;Other

k__Archaea;p__Euryarchaeota