

■ Unassigned:Other:Other:Other
■ K_Archaeap_Euryarchaeota:c_Methanomicrobia.o_Methanocellales
■ K_Archaeap_Euryarchaeota:c_Thermoplasmata.o_Thermoplasmatales
■ K_Archaeap_Miscellaneous Crenarchaeotic Group(SCG):o_Unknown Order
■ K_Archaeap_Thaumarchaeota:Other:Other
■ K_Archaeap_Thaumarchaeota:c_Soil Crenarchaeotic Group(SCG):Other
■ K_Archaeap_Thaumarchaeota:c_Soil Crenarchaeotic Group(SCG):o_Unknown Order
■ K_Archaeap_Thaumarchaeota:c_South African Gold Mine Gp 1(SAGMCG-1):o_Unknown Order
■ K_Archaeap_Thaumarchaeota:c_South African Gold Mine Gp 1(SAGMCG-1):o_Unknown Order
■ K_Archaeap_Thaumarchaeota:c_Territorial group:o_Unknown Order
■ K_Archaeap_Woesearchaeota (DPVE6):c_9_Unknown Order
■ K_Bacteria:Other:Other:Other
■ K_Bacteriap_Acidobacteria:Other:Other
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:Other
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_Unknown Order
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_AT-43-28
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_Acidobacteriales
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_FHC5909
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_Subgroup 11
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_Subgroup 12
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_Subgroup 13
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_Subgroup 15
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_Subgroup 17
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_Subgroup 18
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_Subgroup 19
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_Subgroup 2
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_Subgroup 20
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_Subgroup 21
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_Subgroup 25
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_Subgroup 3
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_Subgroup 4
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_Subgroup 5
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_Subgroup 6
■ K_Bacteriap_Acidobacteria:c_Acidobacteria:o_Subgroup 9
■ K_Bacteriap_Acidobacteria:c_Holophagae:Other
■ K_Bacteriap_Acidobacteria:c_Holophagae:o_Elev-165-816
■ K_Bacteriap_Acidobacteria:c_Holophagae:o_Holophagales
■ K_Bacteriap_Acidobacteria:c_Holophagae:o_Holophagales
■ K_Bacteriap_Acidobacteria:c_Holophagae:o_Subgroup 10
■ K_Bacteriap_Acidobacteria:c_Holophagae:o_Subgroup 7
■ K_Bacteriap_Acidobacteria:c_Holophagae:o_TPD-58
■ K_Bacteriap_Acidobacteria:c_Subgroup 22:o_Unknown Order
■ K_Bacteriap_Acidobacteria:c_Subgroup 26:o_Unknown Order
■ K_Bacteriap_Actinobacteria:Other:Other
■ K_Bacteriap_Actinobacteria:c_Acidimicrobia:o_Acidimicrobiales
■ K_Bacteriap_Actinobacteria:c_Actinobacteria:Other
■ K_Bacteriap_Actinobacteria:c_Actinobacteria:o_Actinomycetales
■ K_Bacteriap_Actinobacteria:c_Actinobacteria:o_Catenulisporales
■ K_Bacteriap_Actinobacteria:c_Actinobacteria:o_Corynebacteriales
■ K_Bacteriap_Actinobacteria:c_Actinobacteria:o_Elev-165-976
■ K_Bacteriap_Actinobacteria:c_Actinobacteria:o_Frankiales
■ K_Bacteriap_Actinobacteria:c_Actinobacteria:o_Glycomycetales
■ K_Bacteriap_Actinobacteria:c_Actinobacteria:o_Kineosporiales
■ K_Bacteriap_Actinobacteria:c_Actinobacteria:o_Micrococcales
■ K_Bacteriap_Actinobacteria:c_Actinobacteria:o_Micromonosporales
■ K_Bacteriap_Actinobacteria:c_Actinobacteria:o_PeM15
■ K_Bacteriap_Actinobacteria:c_Actinobacteria:o_Propionibacteriales
■ K_Bacteriap_Actinobacteria:c_Actinobacteria:o_Pseudonocardiales
■ K_Bacteriap_Actinobacteria:c_Actinobacteria:o_Streptomyetales
■ K_Bacteriap_Actinobacteria:c_Actinobacteria:o_Streptosporangiales
■ K_Bacteriap_Actinobacteria:c_Corobacteriales:o_Corobacteriales
■ K_Bacteriap_Actinobacteria:c_MB-A2-108:o_Unknown Order
■ K_Bacteriap_Actinobacteria:c_Nitrospirales:_Euzeyales
■ K_Bacteriap_Actinobacteria:c_Rubrobacteriales:_Rubrobacteriales
■ K_Bacteriap_Actinobacteria:c_TakashiAC-811:o_Unknown Order
■ K_Bacteriap_Actinobacteria:c_Thermoleophilales:_Thermoleophilales
■ K_Bacteriap_Actinobacteria:c_Thermoleophilales:_Thermoleophilales
■ K_Bacteriap_Aerophobetes:c_9_Unknown Order
■ K_Bacteriap_Armatimonadetes:c_9_Unknown Order
■ K_Bacteriap_Armatimonadetes:c_Armatimonadetes:_Armatimonadales
■ K_Bacteriap_Armatimonadetes:c_Chtionomonadetes:_Chtionomonadales
■ K_Bacteriap_Bacteroidetes:Other:Other
■ K_Bacteriap_Bacteroidetes:c_BSV13:o_Unknown Order
■ K_Bacteriap_Bacteroidetes:c_Bacteroidetes VC2.1_Bac22:o_Unknown Order
■ K_Bacteriap_Bacteroidetes:c_Bacteroidia:o_Bacteroidales
■ K_Bacteriap_Bacteroidetes:c_Bacteroidia:o_Bacteroidia Incertae Sedis
■ K_Bacteriap_Bacteroidetes:c_Cytophagia:o_Cytophagales
■ K_Bacteriap_Bacteroidetes:c_Cytophagia:_Order II
■ K_Bacteriap_Bacteroidetes:c_Flavobacteriales:_Flavobacteriales
■ K_Bacteriap_Bacteroidetes:c_SB-5:o_Unknown Order
■ K_Bacteriap_Bacteroidetes:c_Sphingobacteriales:_Sphingobacteriales
■ K_Bacteriap_Bacteroidetes:c_WCHB1-32:o_Unknown Order
■ K_Bacteriap_Candidate division OP3:c_9_Unknown Order
■ K_Bacteriap_Candidate division SR1:c_9_Unknown Order
■ K_Bacteriap_Candidate division WS6:c_9_Unknown Order
■ K_Bacteriap_Chlamydiae:c_Chlamydiae:_Chlamydiales
■ K_Bacteriap_Chlorobii:_Chlorobiales
■ K_Bacteriap_Chlorobii:_Ignavibacteriales:_Ignavibacteriales
■ K_Bacteriap_Chloroflexi:Other:Other
■ K_Bacteriap_Chloroflexi:c_9_Unknown Order
■ K_Bacteriap_Chloroflexi:c_Anaerolineae:_Anaerolineales
■ K_Bacteriap_Chloroflexi:_Ardentecenia:_Ardenteceniales
■ K_Bacteriap_Chloroflexi:_Caldilineae:_Caldilineales
■ K_Bacteriap_Chloroflexi:_Chloroflexia:_Chloroflexiales
■ K_Bacteriap_Chloroflexi:_Chloroflexia:_Herpetosiphonales
■ K_Bacteriap_Chloroflexi:_Chloroflexia:_Kallotenuales
■ K_Bacteriap_Chloroflexi:_Dehalococcoidia:_Dehalococcoidales
■ K_Bacteriap_Chloroflexi:_Elev-1554:o_Unknown Order
■ K_Bacteriap_Chloroflexi:_Gltt-65-136:o_Unknown Order
■ K_Bacteriap_Chloroflexi:_JG30-KF-CM66:o_Unknown Order
■ K_Bacteriap_Chloroflexi:_JG37-AG-4:o_Unknown Order
■ K_Bacteriap_Chloroflexi:_K04-96:o_Unknown Order
■ K_Bacteriap_Chloroflexi:_Ktedonobacteria:Other
■ K_Bacteriap_Chloroflexi:_Ktedonobacteria:_Ktedonobacteriales
■ K_Bacteriap_Chloroflexi:_Ktedonobacteria:_B10-SB3A
■ K_Bacteriap_Chloroflexi:_Ktedonobacteria:_B12-WMSP1
■ K_Bacteriap_Chloroflexi:_Ktedonobacteria:_C0119
■ K_Bacteriap_Chloroflexi:_Ktedonobacteria:_JG30-KF-A59
■ K_Bacteriap_Chloroflexi:_Ktedonobacteria:_Ktedonobacteriales
■ K_Bacteriap_Chloroflexi:_Ktedonobacteria:_Thermogemmatissporales
■ K_Bacteriap_Chloroflexi:_P2-11E:o_Unknown Order
■ K_Bacteriap_Chloroflexi:_S085:o_Unknown Order
■ K_Bacteriap_Chloroflexi:_SAR202 clade:o_Unknown Order
■ K_Bacteriap_Chloroflexi:_SHA-26:o_Unknown Order
■ K_Bacteriap_Chloroflexi:_TK10:Other
■ K_Bacteriap_Chloroflexi:_TK10:o_Unknown Order
■ K_Bacteriap_Chloroflexi:_TK10:_bacterium Ellin6519
■ K_Bacteriap_Chloroflexi:_TK10:_bacterium Ellin6543
■ K_Bacteriap_Chloroflexi:_Thermomicrobia:_ARXG1722
■ K_Bacteriap_Chloroflexi:_Thermomicrobia:_JG30-KF-CM45
■ K_Bacteriap_Chloroflexi:_Thermomicrobia:_Sphaerobacteriales
■ K_Bacteriap_Cyanobacteria:Other:Other
■ K_Bacteriap_Cyanobacteria:c_Chloroplast:Other
■ K_Bacteriap_Cyanobacteria:c_Chloroplast:o_Unknown Order
■ K_Bacteriap_Cyanobacteria:c_Chloroplast:_Arachis hypogaea var. vulgaris
■ K_Bacteriap_Cyanobacteria:c_Chloroplast:_Cercis gigantha
■ K_Bacteriap_Cyanobacteria:c_Chloroplast:_Lilium perenne
■ K_Bacteriap_Cyanobacteria:c_Chloroplast:_Triticum aestivum (bread wheat)
■ K_Bacteriap_Cyanobacteria:c_Cyanobacteria:Other
■ K_Bacteriap_Cyanobacteria:c_Cyanobacteria:o_Unknown Order
■ K_Bacteriap_Cyanobacteria:c_Cyanobacteria:o_Subsection I
■ K_Bacteriap_Cyanobacteria:c_Cyanobacteria:o_Subsection II
■ K_Bacteriap_Cyanobacteria:c_Cyanobacteria:o_Subsection III
■ K_Bacteriap_Cyanobacteria:c_Cyanobacteria:o_Subsection IV
■ K_Bacteriap_Cyanobacteria:c_ML635/21:o_Unknown Order
■ K_Bacteriap_Cyanobacteria:c_Melainobacteriales:_Obscurbacteriales
■ K_Bacteriap_Cyanobacteria:c_Melainobacteriales:_Vampirobiontales
■ K_Bacteriap_Deferribacteres:c_Deferribacteres:_Deferribacteriales
■ K_Bacteriap_Deinococcus-Thermus:c_Deinococcus:_Deinococcales
■ K_Bacteriap_Elusimicrobia:c_Elusimicrobia:_Elusimicrobiales
■ K_Bacteriap_Elusimicrobia:c_Elusimicrobia:_Lineage I (Endomicrobia)
■ K_Bacteriap_Elusimicrobia:c_Elusimicrobia:_Lineage IIa
■ K_Bacteriap_Elusimicrobia:c_Elusimicrobia:_Lineage IIb
■ K_Bacteriap_Elusimicrobia:c_Elusimicrobia:_Lineage IIc
■ K_Bacteriap_Elusimicrobia:c_Elusimicrobia:_Lineage IV
■ K_Bacteriap_Elusimicrobia:c_Elusimicrobia:_MVP-88
■ K_Bacteriap_Fibrobacteres:c_Fibrobacteriales:_Fibrobacteriales
■ K_Bacteriap_Fibrobacteres:c_Fibrobacteriales:_Fibrobacteria Incertae Sedis
■ K_Bacteriap_Firmicutes:c_Bacilli:_Bacillales
■ K_Bacteriap_Firmicutes:c_Bacilli:_Lactobacillales
■ K_Bacteriap_Firmicutes:c_Clostridia:_Clostridiales
■ K_Bacteriap_Firmicutes:c_Clostridia:_Thermoanaerobacteriales
■ K_Bacteriap_Firmicutes:c_Erysipelotrichia:_Erysipelotrichales
■ K_Bacteriap_Firmicutes:c_Negativicutes:_Selenomonadales
■ K_Bacteriap_Firmicutes:c_OPB54:o_Unknown Order
■ K_Bacteriap_Fusobacteria:c_Fusobacteriales:_Fusobacteriales
■ K_Bacteriap_GAL08:c_9_Unknown Order
■ K_Bacteriap_Gemmatimonadetes:c_Gemmatimonadetes:_AT425-EubC11 terrestrial group
■ K_Bacteriap_Gemmatimonadetes:c_Gemmatimonadetes:_B02-11 terrestrial group
■ K_Bacteriap_Gemmatimonadetes:c_Gemmatimonadetes:_Gemmatimonadales
■ K_Bacteriap_Gemmatimonadetes:c_Gemmatimonadetes:_S0134 terrestrial group
■ K_Bacteriap_Gracilibacteria:c_9_Unknown Order
■ K_Bacteriap_Hydrogenedentes:c_9_Unknown Order
■ K_Bacteriap_JL-ETNP-239:c_9_Unknown Order
■ K_Bacteriap_Latescibacteriac_9_Unknown Order
■ K_Bacteriap_Lentisphaerae:c_Oligosphaeria:_Unknown Order
■ K_Bacteriap_Microgenomates:c_9_Unknown Order
■ K_Bacteriap_Nitrospirae:c_Nitrospirae:_Nitrospirales
■ K_Bacteriap_Omnitrophica:c_NPL-UPA2:o_Unknown Order
■ K_Bacteriap_Parcubacteria:c_9_Unknown Order
■ K_Bacteriap_Planctomycetes:_B07-11:o_Unknown Order
■ K_Bacteriap_Planctomycetes:_C47:o_Unknown Order
■ K_Bacteriap_Planctomycetes:_OM190:o_Unknown Order
■ K_Bacteriap_Planctomycetes:_Phycisphaerae:_CP1a-3 termite group
■ K_Bacteriap_Planctomycetes:_Phycisphaerae:_Phycisphaerales
■ K_Bacteriap_Planctomycetes:_Phycisphaerae:_WD2101 soil group
■ K_Bacteriap_Planctomycetes:_Plal3 lineage:o_Unknown Order
■ K_Bacteriap_Planctomycetes:_Plal4 lineage:o_Unknown Order
■ K_Bacteriap_Planctomycetes:_Planctomycetia:_Planctomycetales
■ K_Bacteriap_Planctomycetes:_SPG12-401-411-872:o_Unknown Order
■ K_Bacteriap_Planctomycetes:_vadinHA49:o_Unknown Order
■ K_Bacteriap_Proteobacteria:Other:Other
■ K_Bacteriap_Proteobacteria:c_ARKICE-90:o_Unknown Order
■ K_Bacteriap_Proteobacteria:c_Alphaproteobacteria:Other
■ K_Bacteriap_Proteobacteria:c_Alphaproteobacteriales:_4-Org1-14
■ K_Bacteriap_Proteobacteria:c_Alphaproteobacteriales:_Caulobacteriales
■ K_Bacteriap_Proteobacteria:c_Alphaproteobacteriales:_DB1-14
■ K_Bacteriap_Proteobacteria:c_Alphaproteobacteriales:_Rhizobiales
■ K_Bacteriap_Proteobacteria:c_Alphaproteobacteriales:_Rhodospirillales
■ K_Bacteriap_Proteobacteria:c_Alphaproteobacteriales:_Rhodospirillales
■ K_Bacteriap_Proteobacteria:c_Alphaproteobacteriales:_Rickettsiales
■ K_Bacteriap_Proteobacteria:c_Alphaproteobacteriales:_Sneathiellales
■ K_Bacteriap_Proteobacteria:c_Alphaproteobacteriales:_Sphingomonadales
■ K_Bacteriap_Proteobacteria:c_Betaproteobacteria:Other
■ K_Bacteriap_Proteobacteria:c_Betaproteobacteriales:_B1-785
■ K_Bacteriap_Proteobacteria:c_Betaproteobacteriales:_Betaproteobacteria Incertae Sedis
■ K_Bacteriap_Proteobacteria:c_Betaproteobacteriales:_Burkholderiales
■ K_Bacteriap_Proteobacteria:c_Betaproteobacteriales:_Hot Creek 32
■ K_Bacteriap_Proteobacteria:c_Betaproteobacteriales:_Hydrogenophilales
■ K_Bacteriap_Proteobacteria:c_Betaproteobacteriales:_Methylophilales
■ K_Bacteriap_Proteobacteria:c_Betaproteobacteriales:_Neisseriales
■ K_Bacteriap_Proteobacteria:c_Betaproteobacteriales:_Nitrosomonadales
■ K_Bacteriap_Proteobacteria:c_Betaproteobacteriales:_Proteobacteriales
■ K_Bacteriap_Proteobacteria:c_Betaproteobacteriales:_Rhodocyclales
■ K_Bacteriap_Proteobacteria:c_Betaproteobacteriales:_SCI-84
■ K_Bacteriap_Proteobacteria:c_Betaproteobacteriales:_TR43-20
■ K_Bacteriap_Proteobacteria:c_Deltaproteobacteria:Other
■ K_Bacteriap_Proteobacteria:c_Deltaproteobacteriales:_43F-1404R
■ K_Bacteriap_Proteobacteria:c_Deltaproteobacteriales:_Bdellovibrionales
■ K_Bacteriap_Proteobacteria:c_Deltaproteobacteriales:_Desulfobacteriales
■ K_Bacteriap_Proteobacteria:c_Deltaproteobacteriales:_Desulfobacteriales
■ K_Bacteriap_Proteobacteria:c_Deltaproteobacteriales:_Desulfurellales
■ K_Bacteriap_Proteobacteria:c_Deltaproteobacteriales:_Desulfurodonadales
■ K_Bacteriap_Proteobacteria:c_Deltaproteobacteriales:_GR-WP33-30
■ K_Bacteriap_Proteobacteria:c_Deltaproteobacteriales:_Myxococcales
■ K_Bacteriap_Proteobacteria:c_Deltaproteobacteriales:_Oligoflexiales
■ K_Bacteriap_Proteobacteria:c_Deltaproteobacteriales:_Proteobacteriales (Marine group B)
■ K_Bacteriap_Proteobacteria:c_Deltaproteobacteriales:_SH1638-Tc1-29
■ K_Bacteriap_Proteobacteria:c_Deltaproteobacteriales:_Syntrophobacteriales
■ K_Bacteriap_Proteobacteria:c_Epsilonproteobacteriales:_Campylobacteriales
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteria:Other
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_1013-28-CG33
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_34P16
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_Aeromonadales
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_B38
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_B072BR169
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_Cellvibrionales
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_Chromatiales
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_EC3
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_Enterobacteriales
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_Gammaproteobacteria Incertae Sedis
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_K189A clade
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_Legionellales
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_Methylococcales
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_NK85
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_Oceanospirillales
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_Order Incertae Sedis
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_PR1103
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_Pasteurellales
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_Pseudomonadales
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_Salinisphaerales
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_Thiotrichales
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_Vibrionales
■ K_Bacteriap_Proteobacteria:c_Gammaproteobacteriales:_Xanthomonadales
■ K_Bacteriap_Proteobacteria:c_TB23:o_Unknown Order
■ K_Bacteriap_Proteobacteria:c_SC3-20:o_Unknown Order
■ K_Bacteriap_Proteobacteria:c_SK259:o_Unknown Order
■ K_Bacteriap_Proteobacteria:c_SPO75OCT0083:o_Unknown Order
■ K_Bacteriap_Proteobacteria:c_TA18:o_Unknown Order
■ K_Bacteriap_Proteobacteria:c_TA18:_Rhizobiales bacterium LX78
■ K_Bacteriap_SHA-109:c_9_Unknown Order
■ K_Bacteriap_SHA211:c_9_Unknown Order
■ K_Bacteriap_Saccharibacteria:Other:Other
■ K_Bacteriap_Saccharibacteriac_9_Unknown Order
■ K_Bacteriap_Saccharibacteriac_Unknown Clade:o_Unknown Order
■ K_Bacteriap_Saccharibacteriac_candidate division TM7 bacterium LY2:o_candidate division TM7 bacterium LY2
■ K_Bacteriap_Spirochaetes:c_Spirochaetes:_Spirochaetales
■ K_Bacteriap_Synergistae:c_Synergistae:_Synergistales
■ K_Bacteriap_TM6:c_9_Unknown Order
■ K_Bacteriap_Tenericutes:c_Mollicutes:_Mollicutes RF9
■ K_Bacteriap_Thermotogae:c_Thermotogae:_Thermotogales
■ K_Bacteriap_Verrucomicrobia:Other:Other
■ K_Bacteriap_Verrucomicrobia:_OPB35 soil group:Other
■ K_Bacteriap_Verrucomicrobia:_OPB35 soil group:o_Unknown Order
■ K_Bacteriap_Verrucomicrobia:_OPB35 soil group:_Unknown Order
■ K_Bacteriap_Verrucomicrobia:_S-BQ2-57 soil group:o_Unknown Order
■ K_Bacteriap_Verrucomicrobia:_Spartobacteriales:_Chthoniobacteriales
■ K_Bacteriap_Verrucomicrobia:_U411:o_Unknown Order
■ K_Bacteriap_Verrucomicrobia:_Verrucomicrobiae Incertae Sedis:_Unknown Order
■ K_Bacteriap_WCHB1-60:c_9_Unknown Order
■ K_Bacteriap_WD272:c_9_Unknown Order