

■ 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_?;o_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_Nitriliruptoria;o_Euzeyrales
■ K_Bacteriap_Actinobacteria;c_Rubrobacteriales;o_Rubrobacteriales
■ K_Bacteriap_Actinobacteria;c_TakashiAC-811;o_Unknown Order
■ K_Bacteriap_Actinobacteria;c_Thermoleophilales;Other
■ K_Bacteriap_Actinobacteria;c_Thermoleophilales;o_Gaellales
■ K_Bacteriap_Actinobacteria;c_Thermoleophilales;o_Solirubrobacteriales
■ K_Bacteriap_Aerophobetes;c_?;o_Unknown Order
■ K_Bacteriap_Armatimonadetes;c_?;o_Unknown Order
■ K_Bacteriap_Armatimonadetes;c_Armatimonadetes;o_Armatimonadales
■ K_Bacteriap_Armatimonadetes;c_Chtionomonadetes;o_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;o_Order II
■ K_Bacteriap_Bacteroidetes;c_Flavobacteriales;o_Flavobacteriales
■ K_Bacteriap_Bacteroidetes;c_SB-5;o_Unknown Order
■ K_Bacteriap_Bacteroidetes;c_Sphingobacteriales;o_Sphingobacteriales
■ K_Bacteriap_Bacteroidetes;c_WCHB1-32;o_Unknown Order
■ K_Bacteriap_Candidate division OP3;c_?;o_Unknown Order
■ K_Bacteriap_Candidate division SR1;c_?;o_Unknown Order
■ K_Bacteriap_Candidate division WS6;c_?;o_Unknown Order
■ K_Bacteriap_Chlamydiae;c_Chlamydiae;o_Chlamydiales
■ K_Bacteriap_Chlorobi;c_Chlorobia;o_Chlorobiales
■ K_Bacteriap_Chlorobi;c_Ignaviibacteriales;o_Ignaviibacteriales
■ K_Bacteriap_Chloroflexi;Other:Other
■ K_Bacteriap_Chloroflexi;c_?;o_Unknown Order
■ K_Bacteriap_Chloroflexi;c_Anaerolineae;o_Anaerolineales
■ K_Bacteriap_Chloroflexi;c_Ardenticatenia;o_Unknown Order
■ K_Bacteriap_Chloroflexi;c_Ardenticatenia;o_Ardenticateniales
■ K_Bacteriap_Chloroflexi;c_Caldilineae;o_Caldilineales
■ K_Bacteriap_Chloroflexi;c_Chloroflexia;o_Chloroflexiales
■ K_Bacteriap_Chloroflexi;c_Chloroflexia;o_Herpetosiphonales
■ K_Bacteriap_Chloroflexi;c_Chloroflexia;o_Kallotenuales
■ K_Bacteriap_Chloroflexi;c_Dehalococcoidia;o_Dehalococcoidales
■ K_Bacteriap_Chloroflexi;c_Elev-1554;o_Unknown Order
■ K_Bacteriap_Chloroflexi;c_GlT-65-136;o_Unknown Order
■ K_Bacteriap_Chloroflexi;c_JG30-KF-CM66;o_Unknown Order
■ K_Bacteriap_Chloroflexi;c_JG37-AG-4;o_Unknown Order
■ K_Bacteriap_Chloroflexi;c_KM4-96;o_Unknown Order
■ K_Bacteriap_Chloroflexi;c_Ktedonobacteria;Other
■ K_Bacteriap_Chloroflexi;c_Ktedonobacteria;o_Unknown Order
■ K_Bacteriap_Chloroflexi;c_Ktedonobacteria;o_B10-SB3A
■ K_Bacteriap_Chloroflexi;c_Ktedonobacteria;o_B12-WMSP1
■ K_Bacteriap_Chloroflexi;c_Ktedonobacteria;o_C0119
■ K_Bacteriap_Chloroflexi;c_Ktedonobacteria;o_JG30-KF-A59
■ K_Bacteriap_Chloroflexi;c_Ktedonobacteria;o_Ktedonobacteriales
■ K_Bacteriap_Chloroflexi;c_Ktedonobacteria;o_Thermogemmatissporales
■ K_Bacteriap_Chloroflexi;c_P2-11E;o_Unknown Order
■ K_Bacteriap_Chloroflexi;c_S085;o_Unknown Order
■ K_Bacteriap_Chloroflexi;c_SAR202 clade;o_Unknown Order
■ K_Bacteriap_Chloroflexi;c_SHA-26;o_Unknown Order
■ K_Bacteriap_Chloroflexi;c_TK10;Other
■ K_Bacteriap_Chloroflexi;c_TK10;o_Unknown Order
■ K_Bacteriap_Chloroflexi;c_TK10;o_Bacterium Ellin6519
■ K_Bacteriap_Chloroflexi;c_TK10;o_Bacterium Ellin6543
■ K_Bacteriap_Chloroflexi;c_Thermomicrobiae;ARXG1722
■ K_Bacteriap_Chloroflexi;c_Thermomicrobiae;JG30-KF-CM45
■ K_Bacteriap_Chloroflexi;c_Thermomicrobiae;o_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;o_Arachis hypogaea var. vulgaris
■ K_Bacteriap_Cyanobacteria;c_Chloroplast;o_Ceris giganea
■ K_Bacteriap_Cyanobacteria;c_Chloroplast;o_Lolium perenne
■ K_Bacteriap_Cyanobacteria;c_Chloroplast;o_Triticum aestivum (bread wheat)
■ K_Bacteriap_Cyanobacteria;c_Cyanobacteria;Other
■ K_Bacteriap_Cyanobacteria;c_Cyanobacteria;o_Unknown Order
■ K_Bacteriap_Cyanobacteria;c_Cyanobacteria;o_SubsectionI
■ K_Bacteriap_Cyanobacteria;c_Cyanobacteria;o_SubsectionII
■ K_Bacteriap_Cyanobacteria;c_Cyanobacteria;o_SubsectionIII
■ K_Bacteriap_Cyanobacteria;c_Cyanobacteria;o_SubsectionIV
■ K_Bacteriap_Cyanobacteria;c_ML635/21;o_Unknown Order
■ K_Bacteriap_Cyanobacteria;c_Melainabacteriales;o_Obscurbacteriales
■ K_Bacteriap_Cyanobacteria;c_Melainabacteriales;o_Vampirovibrionales
■ K_Bacteriap_Deferribacteres;c_Deferribacteres;o_Deferribacteriales
■ K_Bacteriap_Deinococcus-Thermus;c_Deinococcus;o_Deinococcales
■ K_Bacteriap_Elusimicrobia;c_Elusimicrobia;o_Elusimicrobiales
■ K_Bacteriap_Elusimicrobia;c_Elusimicrobia;o_Lineage I (Endomicrobia)
■ K_Bacteriap_Elusimicrobia;c_Elusimicrobia;o_Lineage Ila
■ K_Bacteriap_Elusimicrobia;c_Elusimicrobia;o_Lineage I Ib
■ K_Bacteriap_Elusimicrobia;c_Elusimicrobia;o_Lineage I Ic
■ K_Bacteriap_Elusimicrobia;c_Elusimicrobia;o_Lineage IV
■ K_Bacteriap_Elusimicrobia;c_Elusimicrobia;o_MVP-88
■ K_Bacteriap_Fibrobacteres;c_Fibrobacteria;o_Fibrobacteriales
■ K_Bacteriap_Fibrobacteres;c_Fibrobacteria;o_Fibrobacteria Incertae Sedis
■ K_Bacteriap_Firmicutes;c_Bacilli;o_Bacillales
■ K_Bacteriap_Firmicutes;c_Bacilli;o_Lactobacillales
■ K_Bacteriap_Firmicutes;c_Clostridia;o_Clostridiales
■ K_Bacteriap_Firmicutes;c_Clostridia;o_Thermoanaerobacteriales
■ K_Bacteriap_Firmicutes;c_Erysipelotrichia;o_Erysipelotrichales
■ K_Bacteriap_Firmicutes;c_Negativicutes;o_Selenomonadales
■ K_Bacteriap_Firmicutes;c_OPB54;o_Unknown Order
■ K_Bacteriap_Fusobacteria;c_Fusobacteriales;o_Fusobacteriales
■ K_Bacteriap_GAL08;c_?;o_Unknown Order
■ K_Bacteriap_Gemmatimonadetes;c_Gemmatimonadetes;o_AT425-EubC11 terrestrial group
■ K_Bacteriap_Gemmatimonadetes;c_Gemmatimonadetes;o_B02-11 terrestrial group
■ K_Bacteriap_Gemmatimonadetes;c_Gemmatimonadetes;o_Gemmatimonadales
■ K_Bacteriap_Gemmatimonadetes;c_Gemmatimonadetes;o_S0134 terrestrial group
■ K_Bacteriap_Gracilibacteria;c_?;o_Unknown Order
■ K_Bacteriap_Hydrogenedentes;c_?;o_Unknown Order
■ K_Bacteriap_JL-ETNP-239;c_?;o_Unknown Order
■ K_Bacteriap_Latescibacteriac;c_?;o_Unknown Order
■ K_Bacteriap_Lentisphaerae;c_Oligosphaeria;o_Unknown Order
■ K_Bacteriap_Microgenomates;c_?;o_Unknown Order
■ K_Bacteriap_Nitrospirae;c_Nitrospirae;o_Nitrospirales
■ K_Bacteriap_Omnitrophica;c_NPL-UPA2;o_Unknown Order
■ K_Bacteriap_Parcubacteria;c_?;o_Unknown Order
■ K_Bacteriap_Planctomycetes;c_B07-11;o_Unknown Order
■ K_Bacteriap_Planctomycetes;c_C47;o_Unknown Order
■ K_Bacteriap_Planctomycetes;c_OM190;o_Unknown Order
■ K_Bacteriap_Planctomycetes;c_Phycisphaerae;o_CPla-3 termite group
■ K_Bacteriap_Planctomycetes;c_Phycisphaerae;o_Phycisphaerales
■ K_Bacteriap_Planctomycetes;c_Phycisphaerae;o_WD2101 soil group
■ K_Bacteriap_Planctomycetes;c_Pla3 lineage;o_Unknown Order
■ K_Bacteriap_Planctomycetes;c_Pla4 lineage;o_Unknown Order
■ K_Bacteriap_Planctomycetes;c_Planctomycetia;o_Planctomycetales
■ K_Bacteriap_Planctomycetes;c_SPG12-401-411-872;o_Unknown Order
■ K_Bacteriap_Planctomycetes;c_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_Alphaproteobacteria;o_4-Org1-14
■ K_Bacteriap_Proteobacteria;c_Alphaproteobacteria;o_Caulobacteriales
■ K_Bacteriap_Proteobacteria;c_Alphaproteobacteria;o_DB1-14
■ K_Bacteriap_Proteobacteria;c_Alphaproteobacteria;o_Rhizobiales
■ K_Bacteriap_Proteobacteria;c_Alphaproteobacteria;o_Rhodobacteriales
■ K_Bacteriap_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales
■ K_Bacteriap_Proteobacteria;c_Alphaproteobacteria;o_Rickettsiales
■ K_Bacteriap_Proteobacteria;c_Alphaproteobacteria;o_Sneathiellales
■ K_Bacteriap_Proteobacteria;c_Alphaproteobacteria;o_Sphingomonadales
■ K_Bacteriap_Proteobacteria;c_Betaproteobacteria;Other
■ K_Bacteriap_Proteobacteria;c_Betaproteobacteria;o_B1-785
■ K_Bacteriap_Proteobacteria;c_Betaproteobacteria;o_Betaproteobacteria Incertae Sedis
■ K_Bacteriap_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales
■ K_Bacteriap_Proteobacteria;c_Betaproteobacteria;o_Hot Creek 32
■ K_Bacteriap_Proteobacteria;c_Betaproteobacteria;o_Hydrogenophilales
■ K_Bacteriap_Proteobacteria;c_Betaproteobacteria;o_Methylophilales
■ K_Bacteriap_Proteobacteria;c_Betaproteobacteria;o_Neisseriales
■ K_Bacteriap_Proteobacteria;c_Betaproteobacteria;o_Nitrosomonadales
■ K_Bacteriap_Proteobacteria;c_Betaproteobacteria;o_Prosobranchiales
■ K_Bacteriap_Proteobacteria;c_Betaproteobacteria;o_Rhodocyclales
■ K_Bacteriap_Proteobacteria;c_Betaproteobacteria;o_SCI-84
■ K_Bacteriap_Proteobacteria;c_Betaproteobacteria;o_TRA3-20
■ K_Bacteriap_Proteobacteria;c_Deltaproteobacteria;Other
■ K_Bacteriap_Proteobacteria;c_Deltaproteobacteria;o_43F-1404R
■ K_Bacteriap_Proteobacteria;c_Deltaproteobacteria;o_Bdellovibrionales
■ K_Bacteriap_Proteobacteria;c_Deltaproteobacteria;o_Desulfobacteriales
■ K_Bacteriap_Proteobacteria;c_Deltaproteobacteria;o_Desulfobacteriales
■ K_Bacteriap_Proteobacteria;c_Deltaproteobacteria;o_Desulfurellales
■ K_Bacteriap_Proteobacteria;c_Deltaproteobacteria;o_Desulfuromonadales
■ K_Bacteriap_Proteobacteria;c_Deltaproteobacteria;o_GR-WP53-30
■ K_Bacteriap_Proteobacteria;c_Deltaproteobacteria;o_Myxococcales
■ K_Bacteriap_Proteobacteria;c_Deltaproteobacteria;o_Oligoflexales
■ K_Bacteriap_Proteobacteria;c_Deltaproteobacteria;o_Prosobranchiales
■ K_Bacteriap_Proteobacteria;c_Deltaproteobacteria;o_Siniphilales
■ K_Bacteriap_Proteobacteria;c_Deltaproteobacteria;o_Siniphilales
■ K_Bacteriap_Proteobacteria;c_Deltaproteobacteria;o_Syntrophobacteriales
■ K_Bacteriap_Proteobacteria;c_Epsilonproteobacteria;o_Campylobacteriales
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;Other
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_Unknown Order
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_1013-28-CG33
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_34P16
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_Aeromonadales
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_B38
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_BD72BR169
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_Cellvibrionales
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_Chromatiales
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_EC3
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_Enterobacteriales
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_Gammaproteobacteria Incertae Sedis
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_K189A clade
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_Legionellales
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_Methylococcales
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_NK85
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_Oceanospirillales
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_Order Incertae Sedis
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_PPR1063
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_Pasteurellales
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_Pseudomonadales
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_Salinisphaerales
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_Thiotrichales
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_Vibrionales
■ K_Bacteriap_Proteobacteria;c_Gammaproteobacteria;o_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_SPO7SOCT00n83;o_Unknown Order
■ K_Bacteriap_Proteobacteria;c_TAJ18;o_Unknown Order
■ K_Bacteriap_Proteobacteria;c_TA18;o_Rhizobiales bacterium LX78
■ K_Bacteriap_SHA-109;c_?;o_Unknown Order
■ K_Bacteriap_SHA211;c_?;o_Unknown Order
■ K_Bacteriap_Saccharibacteria;Other:Other
■ K_Bacteriap_Saccharibacteria;c_?;o_Unknown Order
■ K_Bacteriap_Saccharibacteria;c_Unknown Class;o_Unknown Order
■ K_Bacteriap_Saccharibacteria;c_candidate division TM7 bacterium LY2;o_candidate division TM7 bacterium LY2
■ K_Bacteriap_Spirochaetes;c_Spirochaetes;o_Spirochaetales
■ K_Bacteriap_Synergistae;c_Synergistae;o_Synergistales
■ K_Bacteriap_TM6;c_?;o_Unknown Order
■ K_Bacteriap_Tenericutes;c_Mollicutes;o_Mollicutes RF9
■ K_Bacteriap_Thermotogae;c_Thermotogae;o_Thermotogales
■ K_Bacteriap_Verrucomicrobia;Other:Other
■ K_Bacteriap_Verrucomicrobia;c_OPB35 soil group;Other
■ K_Bacteriap_Verrucomicrobia;c_OPB35 soil group;o_Unknown Order
■ K_Bacteriap_Verrucomicrobia;c_OPB35 soil group;o_Unknown Order
■ K_Bacteriap_Verrucomicrobia;c_S-BQ2-57 soil group;o_Unknown Order
■ K_Bacteriap_Verrucomicrobia;c_Spartobacteria;o_Chthoniobacteriales
■ K_Bacteriap_Verrucomicrobia;c_UA11;o_Unknown Order
■ K_Bacteriap_Verrucomicrobia;c_Verrucomicrobiae Incertae Sedis;o_Unknown Order
■ K_Bacteriap_Verrucomicrobia;c_Verrucomicrobiae;o_Verrucomicrobiales
■ K_Bacteriap_WCHB1-60;c_?;o_Unknown Order
■ K_Bacteriap_WD272;c_?;o_Unknown Order