

- Unassigned;Other;Other
- k__Archaea;p__Crenarchaeota;c__MBGA
- k__Archaea;p__Crenarchaeota;c__MBGB
- k__Archaea;p__Crenarchaeota;c__MCG
- k__Archaea;p__Crenarchaeota;c__Thaumarchaeota
- k__Archaea;p__Euryarchaeota;c__ANME-1
- k__Archaea;p__Euryarchaeota;c__Halobacteria
- k__Archaea;p__Euryarchaeota;c__Methanobacteria
- k__Archaea;p__Euryarchaeota;c__Methanomicrobia
- k__Archaea;p__Euryarchaeota;c__Thermoplasmata
- k__Archaea;p__[Parvarchaeota];c__[Parvarchaea]
- k__Bacteria;Other;Other
- k__Bacteria;p__;c__
- k__Bacteria;p__AC1;c__HDBW-WB69
- k__Bacteria;p__AD3;c__ABS-6
- k__Bacteria;p__AD3;c__JG37-AG-4
- k__Bacteria;p__Acidobacteria;c__
- k__Bacteria;p__Acidobacteria;c__AT-s2-57
- k__Bacteria;p__Acidobacteria;c__AT-s54
- k__Bacteria;p__Acidobacteria;c__Acidobacteria-5
- k__Bacteria;p__Acidobacteria;c__Acidobacteria-6
- k__Bacteria;p__Acidobacteria;c__Acidobacteriia
- k__Bacteria;p__Acidobacteria;c__BPC102
- k__Bacteria;p__Acidobacteria;c__DA052
- k__Bacteria;p__Acidobacteria;c__EC1113
- k__Bacteria;p__Acidobacteria;c__Holophagae
- k__Bacteria;p__Acidobacteria;c__OS-K
- k__Bacteria;p__Acidobacteria;c__PAUC37f
- k__Bacteria;p__Acidobacteria;c__RB25
- k__Bacteria;p__Acidobacteria;c__S035
- k__Bacteria;p__Acidobacteria;c__Solibacteres
- k__Bacteria;p__Acidobacteria;c__Sva0725
- k__Bacteria;p__Acidobacteria;c__TM1
- k__Bacteria;p__Acidobacteria;c__[Chloracidobacteria]
- k__Bacteria;p__Acidobacteria;c__iii1-8
- k__Bacteria;p__Actinobacteria;Other
- k__Bacteria;p__Actinobacteria;c__
- k__Bacteria;p__Actinobacteria;c__
- k__Bacteria;p__Actinobacteria;c__Acidimicrobia
- k__Bacteria;p__Actinobacteria;c__Actinobacteria
- k__Bacteria;p__Actinobacteria;c__Coriobacteriia
- k__Bacteria;p__Actinobacteria;c__MB-A2-108
- k__Bacteria;p__Actinobacteria;c__Nitriliruptoria
- k__Bacteria;p__Actinobacteria;c__OPB41
- k__Bacteria;p__Actinobacteria;c__Rubrobacteria
- k__Bacteria;p__Actinobacteria;c__Thermoleophilia
- k__Bacteria;p__AncK6;c__
- k__Bacteria;p__Aquificae;c__Aquificae
- k__Bacteria;p__Armatimonadetes;c__0319-6E2
- k__Bacteria;p__Armatimonadetes;c__Armatimonadia
- k__Bacteria;p__Armatimonadetes;c__Chthonomonadetes
- k__Bacteria;p__Armatimonadetes;c__OPB50
- k__Bacteria;p__Armatimonadetes;c__SHA-37
- k__Bacteria;p__Armatimonadetes;c__SJA-176
- k__Bacteria;p__Armatimonadetes;c__[Fimbriimonadia]
- k__Bacteria;p__BHI80-139;c__
- k__Bacteria;p__BRC1;c__PRR-11
- k__Bacteria;p__Bacteroidetes;Other
- k__Bacteria;p__Bacteroidetes;c__At12OctB3
- k__Bacteria;p__Bacteroidetes;c__BME43
- k__Bacteria;p__Bacteroidetes;c__Bacteroidia
- k__Bacteria;p__Bacteroidetes;c__Cytophagia
- k__Bacteria;p__Bacteroidetes;c__Flavobacteriia
- k__Bacteria;p__Bacteroidetes;c__SM1A07
- k__Bacteria;p__Bacteroidetes;c__Sphingobacteriia
- k__Bacteria;p__Bacteroidetes;c__VC2_1_Bac22
- k__Bacteria;p__Bacteroidetes;c__[Rhodothermi]
- k__Bacteria;p__Bacteroidetes;c__[Saprospirae]
- k__Bacteria;p__Caldiseria;c__WCHB1-03
- k__Bacteria;p__Caldithrix;c__Caldithrixae
- k__Bacteria;p__Chlamydiae;c__Chlamydia
- k__Bacteria;p__Chlorobi;c__
- k__Bacteria;p__Chlorobi;c__BSV26
- k__Bacteria;p__Chlorobi;c__Ignavibacteria
- k__Bacteria;p__Chlorobi;c__OPB56
- k__Bacteria;p__Chlorobi;c__SJA-28
- k__Bacteria;p__Chloroflexi;Other
- k__Bacteria;p__Chloroflexi;c__
- k__Bacteria;p__Chloroflexi;c__Anaerolineae
- k__Bacteria;p__Chloroflexi;c__C0119
- k__Bacteria;p__Chloroflexi;c__Chloroflexi
- k__Bacteria;p__Chloroflexi;c__Dehalococcoidetes
- k__Bacteria;p__Chloroflexi;c__Ellin6529
- k__Bacteria;p__Chloroflexi;c__Gitt-GS-136
- k__Bacteria;p__Chloroflexi;c__Ktedonobacteria
- k__Bacteria;p__Chloroflexi;c__P2-11E
- k__Bacteria;p__Chloroflexi;c__S085
- k__Bacteria;p__Chloroflexi;c__SAR202
- k__Bacteria;p__Chloroflexi;c__SHA-26
- k__Bacteria;p__Chloroflexi;c__TK10
- k__Bacteria;p__Chloroflexi;c__TK17
- k__Bacteria;p__Chloroflexi;c__Thermomicrobia
- k__Bacteria;p__Chloroflexi;c__[Thermobacula]
- k__Bacteria;p__Cyanobacteria;Other
- k__Bacteria;p__Cyanobacteria;c__
- k__Bacteria;p__Cyanobacteria;c__4C0d-2
- k__Bacteria;p__Cyanobacteria;c__Chloroplast
- k__Bacteria;p__Cyanobacteria;c__Gloeobacteriophycideae
- k__Bacteria;p__Cyanobacteria;c__ML635J-21
- k__Bacteria;p__Cyanobacteria;c__Nostocophycideae
- k__Bacteria;p__Cyanobacteria;c__Oscillatoriohycideae
- k__Bacteria;p__Cyanobacteria;c__Synechococcophycideae
- k__Bacteria;p__Deferribacteres;c__Deferribacteres
- k__Bacteria;p__Elusimicrobia;c__
- k__Bacteria;p__Elusimicrobia;c__Elusimicrobia
- k__Bacteria;p__Elusimicrobia;c__Endomicrobia
- k__Bacteria;p__FBP;c__
- k__Bacteria;p__FCPU426;c__
- k__Bacteria;p__Fibrobacteres;c__
- k__Bacteria;p__Fibrobacteres;c__B5-096
- k__Bacteria;p__Fibrobacteres;c__Fibrobacteria
- k__Bacteria;p__Fibrobacteres;c__SBZC_2415
- k__Bacteria;p__Fibrobacteres;c__TG3
- k__Bacteria;p__Firmicutes;Other
- k__Bacteria;p__Firmicutes;c__Bacilli
- k__Bacteria;p__Firmicutes;c__Clostridia
- k__Bacteria;p__Firmicutes;c__Erysipelotrichi
- k__Bacteria;p__Firmicutes;c__OPB54
- k__Bacteria;p__Fusobacteria;c__Fusobacteriia
- k__Bacteria;p__GN02;c__
- k__Bacteria;p__GN02;c__BB34
- k__Bacteria;p__GN02;c__BD1-5
- k__Bacteria;p__GN02;c__GKS2-174
- k__Bacteria;p__GN02;c__GN07
- k__Bacteria;p__GN04;c__
- k__Bacteria;p__GN04;c__GN15
- k__Bacteria;p__GN04;c__MSB-5A5
- k__Bacteria;p__GOUTA4;c__
- k__Bacteria;p__Gemmatimonadetes;c__
- k__Bacteria;p__Gemmatimonadetes;c__Gemm-1
- k__Bacteria;p__Gemmatimonadetes;c__Gemm-2
- k__Bacteria;p__Gemmatimonadetes;c__Gemm-3
- k__Bacteria;p__Gemmatimonadetes;c__Gemm-4
- k__Bacteria;p__Gemmatimonadetes;c__Gemm-5
- k__Bacteria;p__Gemmatimonadetes;c__Gemmatimonadetes
- k__Bacteria;p__Kazan-3B-28;c__
- k__Bacteria;p__LD1;c__
- k__Bacteria;p__Lentisphaerae;c__[Lentisphaeria]
- k__Bacteria;p__MVP-21;c__
- k__Bacteria;p__NC10;c__12-24
- k__Bacteria;p__NC10;c__wb1-A12
- k__Bacteria;p__NKB19;c__TSBW08
- k__Bacteria;p__Nitrospirae;c__Nitrospira
- k__Bacteria;p__OC31;c__
- k__Bacteria;p__OD1;c__
- k__Bacteria;p__OD1;c__ABY1
- k__Bacteria;p__OD1;c__Mb-NB09
- k__Bacteria;p__OD1;c__SM2F11
- k__Bacteria;p__OD1;c__ZB2
- k__Bacteria;p__OP11;c__
- k__Bacteria;p__OP11;c__OP11-2
- k__Bacteria;p__OP11;c__OP11-3
- k__Bacteria;p__OP11;c__OP11-4
- k__Bacteria;p__OP11;c__WCHB1-64
- k__Bacteria;p__OP3;c__
- k__Bacteria;p__OP3;c__BD4-9
- k__Bacteria;p__OP3;c__PBS-25
- k__Bacteria;p__OP3;c__kol11
- k__Bacteria;p__OP8;c__OP8_1
- k__Bacteria;p__OP8;c__SAW1_B6
- k__Bacteria;p__OP9;c__JS1
- k__Bacteria;p__PAUC34f;c__
- k__Bacteria;p__Planctomycetes;c__
- k__Bacteria;p__Planctomycetes;c__028H05-P-BN-P5
- k__Bacteria;p__Planctomycetes;c__BD7-11
- k__Bacteria;p__Planctomycetes;c__C6
- k__Bacteria;p__Planctomycetes;c__ODP123
- k__Bacteria;p__Planctomycetes;c__OM190
- k__Bacteria;p__Planctomycetes;c__Phycisphaerae
- k__Bacteria;p__Planctomycetes;c__Pla4
- k__Bacteria;p__Planctomycetes;c__Planctomycetia
- k__Bacteria;p__Planctomycetes;c__SBZP_4958
- k__Bacteria;p__Planctomycetes;c__[Brocadiae]
- k__Bacteria;p__Planctomycetes;c__vadinHA49
- k__Bacteria;p__Proteobacteria;Other
- k__Bacteria;p__Proteobacteria;c__
- k__Bacteria;p__Proteobacteria;c__Alphaproteobacteria
- k__Bacteria;p__Proteobacteria;c__Betaproteobacteria
- k__Bacteria;p__Proteobacteria;c__Deltaproteobacteria
- k__Bacteria;p__Proteobacteria;c__Epsilonproteobacteria
- k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria
- k__Bacteria;p__Proteobacteria;c__TA18
- k__Bacteria;p__SAR406;c__AB16
- k__Bacteria;p__SBR1093;c__
- k__Bacteria;p__SBR1093;c__A712011
- k__Bacteria;p__SC4;c__
- k__Bacteria;p__SR1;c__
- k__Bacteria;p__Spirochaetes;c__MVP-15
- k__Bacteria;p__Spirochaetes;c__Spirochaetes
- k__Bacteria;p__Spirochaetes;c__[Brachyspirae]
- k__Bacteria;p__Spirochaetes;c__[Brevinematae]
- k__Bacteria;p__Spirochaetes;c__[Leptospirae]
- k__Bacteria;p__Synergistetes;c__Synergistia
- k__Bacteria;p__TM6;c__F38
- k__Bacteria;p__TM6;c__SBRH58
- k__Bacteria;p__TM6;c__SJA-4
- k__Bacteria;p__TM7;c__
- k__Bacteria;p__TM7;c__MJK10
- k__Bacteria;p__TM7;c__SC3
- k__Bacteria;p__TM7;c__TM7-1
- k__Bacteria;p__TM7;c__TM7-3
- k__Bacteria;p__Tenericutes;c__CK-1C4-19
- k__Bacteria;p__Tenericutes;c__Mollicutes
- k__Bacteria;p__Tenericutes;c__RF3
- k__Bacteria;p__Thermotogae;c__Thermotogae
- k__Bacteria;p__Verrucomicrobia;c__Opitutae
- k__Bacteria;p__Verrucomicrobia;c__Verruco-5
- k__Bacteria;p__Verrucomicrobia;c__Verrucomicrobiae
- k__Bacteria;p__Verrucomicrobia;c__[Methylacidiphilae]
- k__Bacteria;p__Verrucomicrobia;c__[Pedosphaerae]
- k__Bacteria;p__Verrucomicrobia;c__[Spartobacteria]
- k__Bacteria;p__WPS-2;c__
- k__Bacteria;p__WS1;c__
- k__Bacteria;p__WS2;c__Kazan-3B-09
- k__Bacteria;p__WS2;c__PRA-109
- k__Bacteria;p__WS3;c__SHR-12
- k__Bacteria;p__WS4;c__
- k__Bacteria;p__WS5;c__
- k__Bacteria;p__WWE1;c__MSBL2
- k__Bacteria;p__WWE1;c__[Cloacamonae]
- k__Bacteria;p__ZB3;c__
- k__Bacteria;p__ZB3;c__BS119
- k__Bacteria;p__ZB3;c__KSb1
- k__Bacteria;p__[Thermi];c__Deinococci