

Reference_U03403.1_Microcystis_aeruginosa_NIES89_16S_rRNA_gene
 Oxic_PS_U03403.1_Microcystis_aeruginosa_NIES89_16S_rRNA_gene
 Oxic_PS_AJ635429.1_Microcystis_aeruginosa_1BB38S07_partial_16S_rRNA_gene
 Oxic_PS_AJ635430.1_Microcystis_aeruginosa_0BB35S02_partial_16S_rRNA_gene
 Reference_U03402.1_Microcystis_aeruginosa_PCC7806_16S_rRNA_gene
 Oxic_PS_U03402.1_Microcystis_aeruginosa_PCC7806_16S_rRNA_gene
 Reference_U40338.1_Microcystis_aeruginosa_PCC7005_16S_rRNA_gene(2)
 Reference_U40338.1_Microcystis_aeruginosa_PCC7005_16S_rRNA_gene
 Oxic_PS_U40338.1_Microcystis_aeruginosa_PCC7005_16S_rRNA_gene
 Oxic_PS_AJ635431.1_Microcystis_aeruginosa_0BF29S01_partial_16S_rRNA_gene
 Reference_AJ635432.1_Microcystis_aeruginosa_0BF29S03_partial_16S_rRNA_gene
 Reference_AJ635431.1_Microcystis_aeruginosa_0BF29S01_partial_16S_rRNA_gene(2)
 Reference_AJ635431.1_Microcystis_aeruginosa_0BF29S01_partial_16S_rRNA_gene
 Oxic_PS_AJ635432.1_Microcystis_aeruginosa_0BF29S03_partial_16S_rRNA_gene
 Reference_AJ133156.1_Anabaena_sp._90_partial_16S_rRNA_gene_strain_90
 Oxic_PS_AJ133156.1_Anabaena_sp._90_partial_16S_rRNA_gene_strain_90
 Oxic_PS_HE975014.1_Anabaena_cylindrica_PCC_7122_partial_16S_rRNA_gene_strain_CCAP_1403/2B
 Oxic_PS_AJ488603.1_Anabaena_azotica_partial_16S_rRNA_gene
 Reference_AM236076.1_Planktothrix_pseudagardhii_partial_16S_rRNA_gene_isolate_VR1
 Oxic_PS_AM236076.1_Planktothrix_pseudagardhii_partial_16S_rRNA_gene_isolate_VR1
 Oxic_PS_AJ132252.1_Planktothrix_rubescens_16S_rRNA_gene_partial_strain_BC-Pla_9316
 Oxic_PS_AJ544068.1_Planktothrix_rubescens_partial_16S_rRNA_gene_strain_PMC11002
 Reference_AJ133185.1_Planktothrix_sp._2_partial_16S_rRNA_gene_strain_2
 Oxic_PS_AJ133185.1_Planktothrix_sp._2_partial_16S_rRNA_gene_strain_2
 Reference_AJ635435.1_Planktothrix_sp._1LT27S08_partial_16S_rRNA_gene
 Oxic_PS_AJ635435.1_Planktothrix_sp._1LT27S08_partial_16S_rRNA_gene
 Anoxic PS AB234548.1_Uncultured_Chloroflexi_bacterium_gene_for_16S_rRNA_partial_sequence_clone:MgMjR-045
 Reference_KR088538.1.1521_BacteriaActinobacteriaActinobacteriaMicrococcalesMicrobacteriaceaeLeifsoniaLeifsonia_sp._D24
 Reference_JOOI01000002.388919.390619_BacteriaFirmicutesClostridiaThermoanaerobacteralesThermoanaerobacterThermoanaerobacter_sp._YS13
 Reference_AEYS01000010.10484.12283_BacteriaFirmicutesClostridiaThermoanaerobacteralesThermoanaerobacteraceaeThermoanaerobacterThermoanaerobacter_ethanolicus_JW_200
 Reference_LC127108.1.1643_BacteriaFirmicutesClostridiaThermoanaerobacteralesFamily_IIIThermoanaerobacteriumThermoanaerobacterium_sp.
 Reference_AF321086.1.1725_BacteriaFirmicutesClostridiaThermoanaerobacteralesThermoanaerobacteraceaeGelriaGelriaGlutamica
 Reference_CP000860.1442921.1444612_BacteriaFirmicutesClostridiaClostridia_Incertae_SedisUnknown_FamilyCandidatus_DesulforudisCandidatus_Desulforudis_audaxviator_MP104C
 Reference_JQID01000098.234.2000_BacteriaFirmicutesClostridiaClostridialesPeptococcaceaeDesulfohalobiumDesulfohalobium_sp._Tol-M
 Anoxic PS LN831324.1_Heliobacterium_gestii_partial_16S_rRNA_gene_strain_DSM_11169_clone_10
 Anoxic PS LN831323.1_Heliobacterium_gestii_partial_16S_rRNA_gene_strain_DSM_11169_clone_9
 Anoxic PS HE582751.1_Heliophilum_fasciatum_partial_16S_rRNA_gene_type_strain_DSM_11170T_clone_1
 Anoxic PS HE582753.1_Heliophilum_fasciatum_partial_16S_rRNA_gene_type_strain_DSM_11170T_clone_3
 Anoxic PS HE582752.1_Heliophilum_fasciatum_partial_16S_rRNA_gene_type_strain_DSM_11170T_clone_2
 Anoxic PS NR_137410.1_Terriglobus_albidus_strain_Ac_26_B10_16S_ribosomal_RNA_partial_sequence
 Anoxic PS NR_133877.1_Terriglobus_tenax_strain_DRP_35_16S_ribosomal_RNA_partial_sequence
 Anoxic PS NR_147738.1_Occallatibacter_riparius_strain_277_16S_ribosomal_RNA_partial_sequence
 Anoxic PS NR_147737.1_Occallatibacter_savannae_strain_A2-1c_16S_ribosomal_RNA_partial_sequence
 Reference_CP012147.1239597.1241161_BacteriaNitrospiraeNitrospiraeNitrospiralesNitrospiraceaeLeptospirillumLeptospirillum_sp._Group_II_CF-1
 Reference_LWDU01036823.11333.12835_BacteriaVerrucomicrobiaOPB35_soil_group_hydrothermal_vent_metagenome
 Reference_KM210554.1.1546_BacteriaVerrucomicrobiaVerrucomicrobia_Incertae_SedisUnknown_OrderUnknown_FamilyCandidatus_MethylacidiphilumMethylacidimicrobium_tartarophyllum
 Reference_CCPS01000022.154.1916_BacteriaProteobacteriaGammaproteobacteriaEnterobacterialesEnterobacteriaceaeEscherichia-ShigellaEscherichia_coli
 Reference_AOQJ01000007.3335.4955_BacteriaProteobacteriaGammaproteobacteriaEnterobacterialesEnterobacteriaceaeEscherichia-ShigellaEscherichia_coli_TOP293-4
 Reference_JQ924831.1.1625_BacteriaProteobacteriaGammaproteobacteriaEnterobacterialesEnterobacteriaceaePhotobacteriumPhotobacterium_luminescens
 Reference_EF636184.1.917_BacteriaProteobacteriaGammaproteobacteriaChromatialesChromatiaceaeRheinheimeraChromatiales_bacterium_TP178
 Anoxic PS NR_118133.1_Halothiobacillus_neapolitanus_strain_NCIMB_8539_16S_ribosomal_RNA_gene_partial_sequence
 Anoxic PS NR_104929.1_Halothiobacillus_neapolitanus_strain_Parker_strain_X_16S_ribosomal_RNA_gene_complete_sequence
 Anoxic PS NR_117125.1_Thioalkalivibrio_versutus_strain_DSM_13738_16S_ribosomal_RNA_gene_partial_sequence
 Anoxic PS NR_114774.1_Nitrosococcus_halophilus_strain_Nc_4_16S_ribosomal_RNA_gene_partial_sequence
 Anoxic PS NR_026245.1_Rhabdochromatium_marinum_strain_DSM_5261_16S_ribosomal_RNA_gene_partial_sequence
 Anoxic PS NR_113390.1_Halorhodospira_neutrophila_strain_JCM_17835_16S_ribosomal_RNA_gene_partial_sequence
 Anoxic PS NR_104779.1_Halorhodospira_halochloris_strain_BN_9850_16S_ribosomal_RNA_gene_partial_sequence
 Anoxic PS NR_104763.1_Rhodovibrio_sodomensis_strain_DSI_16S_ribosomal_RNA_gene_partial_sequence
 Reference_AF513443.1.1398_BacteriaProteobacteriaAlphaproteobacteriaRhodospirillalesRhodospirillaceaeRhodospirillumRhodospirillum_sp._LA45
 Reference_EU704115.1.1487_BacteriaProteobacteriaAlphaproteobacteriaRhodospirillalesRhodospirillaceaeThalassospiraRhodospirillaceae_bacterium_EZ54
 Reference_DQ270717.1.1333_BacteriaProteobacteriaAlphaproteobacteriaRhodospirillalesRhodospirillaceaeThalassospiraRhodospirillaceae_bacterium_B-2164
 Reference_AB371589.1.1416_BacteriaProteobacteriaAlphaproteobacteriaRhodospirillalesRhodospirillaceaeRhodospirillumRhodospirillum_sp._03PA-Blue-76
 Anoxic PS NR_104755.1_Phaeospirillum_molischianum_strain_DSM_120_16S_ribosomal_RNA_gene_partial_sequence
 Anoxic PS NR_117477.1_Azospirillum_amazonense_strain_DSM_2787_16S_ribosomal_RNA_gene_partial_sequence
 Anoxic PS NR_104981.1_Azospirillum_amazonense_strain_Am_14_16S_ribosomal_RNA_gene_partial_sequence
 Anoxic PS NR_117480.1_Azospirillum_irakense_strain_DSM_11586_16S_ribosomal_RNA_gene_partial_sequence
 Anoxic PS NR_117870.1_Azospirillum_canadense_strain_LMG_23617_16S_ribosomal_RNA_gene_partial_sequence
 Anoxic PS NR_117478.1_Azospirillum_brasiliense_strain_DSM_1690_16S_ribosomal_RNA_gene_partial_sequence
 Anoxic PS NR_117479.1_Azospirillum_halopraeferens_strain_DSM_3675_16S_ribosomal_RNA_gene_partial_sequence
 Anoxic PS NR_117482.1_Azospirillum_oryzae_strain_JCM_21588_16S_ribosomal_RNA_gene_partial_sequence
 Anoxic PS NR_117481.1_Azospirillum_lipoferum_strain_DSM_1691_16S_ribosomal_RNA_gene_partial_sequence
 Anoxic PS NR_104865.1_Chlorobaculum_limnaeum_strain_DSM_1677_16S_ribosomal_RNA_gene_partial_sequence
 Anoxic PS AH001601.2_Chloroflexus_aurantiacus_strain_Y400-fl_16S_ribosomal_RNA_(16S_rRNA)_complete_sequence
 GU227339.1_Uncultured_marine_archaeon_clone_BArc12_16S_ribosomal_RNA_gene_partial_sequence