

D_0_ArchaeaD_1_EuryarchaeotaD_2_Methanobacteria.D_3_Methanobacteriales.D_4_MethanobacteriaceaeD_5_Methanobacterium

D_0_BacteriaD_1_ActinobacteriaD_2_ActinobacteriaD_3_SoliribacteralesD_4_SoliribacteriaceaeD_5_Candidatus Soliribacter

D_0_BacteriaD_1_ActinobacteriaD_2_ActinobacteriaD_3_CorynebacterialesD_4_CorynebacteriaceaeD_5_Corynebacterium 1

D_0_BacteriaD_1_ActinobacteriaD_2_ActinobacteriaD_3_MicrococcalesD_4_BogoriellaceaeD_5_Georgia

D_0_BacteriaD_1_ActinobacteriaD_2_ActinobacteriaD_3_MicrococcalesD_4_CellulomonadaceaeD_5_Actinozeta

D_0_BacteriaD_1_ActinobacteriaD_2_ActinobacteriaD_3_MicrococcalesD_4_CellulomonadaceaeD_5_Cellulomonas

D_0_BacteriaD_1_ActinobacteriaD_2_ActinobacteriaD_3_MicrococcalesD_4_DemequiniaceaeOther

D_0_BacteriaD_1_ActinobacteriaD_2_ActinobacteriaD_3_MicrococcalesD_4_MicrobacteriaceaeAmbiguous taxa

D_0_BacteriaD_1_ActinobacteriaD_2_ActinobacteriaD_3_MicrococcalesD_4_MicrobacteriaceaeD_5_Pseudocylindroclavibacter

D_0_BacteriaD_1_ActinobacteriaD_2_ActinobacteriaD_3_MicrococcalesD_4_MicrococcaeaeD_5_Nesterenkonia

D_0_BacteriaD_1_ActinobacteriaD_2_CorioribacteralesD_3_PseudonocardialesD_4_PseudonocardiaecaeD_5_Pseudonocardia

D_0_BacteriaD_1_ActinobacteriaD_2_CorioribacteralesD_3_CorioribacteralesD_4_CorioribacteriaceaeD_5_Enterorhabdus

D_0_BacteriaD_1_ActinobacteriaD_2_CorioribacteralesD_3_CorioribacteralesD_4_CorioribacteriaceaeD_5_Parvibacter

D_0_BacteriaD_1_ActinobacteriaD_2_CorioribacteralesD_3_CorioribacteralesD_4_CorioribacteriaceaeD_5_uncultured

D_0_BacteriaD_1_ActinobacteriaD_2_CorioribacteralesD_3_CorioribacteralesD_4_CorioribacteriaceaeOther

D_0_BacteriaD_1_ActinobacteriaD_2_ThermopileophilaD_3_GailleriesD_4_unculturedD_5_uncultured bacterium

D_0_BacteriaD_1_ActinobacteriaD_2_ThermopileophilaD_3_SoliribacteralesD_4_Elen-165132D_5_uncultured microorganism

D_0_BacteriaD_1_BacteroidetesD_2_BacteroidiaD_3_BacteroidalesD_4_BacteroidaceaeD_5_Bacteroides

D_0_BacteriaD_1_BacteroidetesD_2_BacteroidiaD_3_BacteroidalesD_4_Bacteroidales S24-7 groupD_5_mouse gut metagenome

D_0_BacteriaD_1_BacteroidetesD_2_BacteroidiaD_3_BacteroidalesD_4_Bacteroidales S24-7 groupD_5_uncultured bacterium

D_0_BacteriaD_1_BacteroidetesD_2_BacteroidiaD_3_BacteroidalesD_4_Bacteroidales S24-7 groupOther

D_0_BacteriaD_1_BacteroidetesD_2_BacteroidiaD_3_BacteroidalesD_4_PorphyrionomadaeaeD_5_Dysgonomonas

D_0_BacteriaD_1_BacteroidetesD_2_BacteroidiaD_3_BacteroidalesD_4_PorphyrionomadaeaeD_5_Parabacteroides

D_0_BacteriaD_1_BacteroidetesD_2_BacteroidiaD_3_BacteroidalesD_4_PorphyrionomadaeaeD_5_uncultured

D_0_BacteriaD_1_BacteroidetesD_2_BacteroidiaD_3_BacteroidalesD_4_PrevotellaceaeD_5_Prevotella 9

D_0_BacteriaD_1_BacteroidetesD_2_BacteroidiaD_3_BacteroidalesD_4_PrevotellaceaeD_5_Prevotellaceae NK0831 group

D_0_BacteriaD_1_BacteroidetesD_2_BacteroidiaD_3_BacteroidalesD_4_PrevotellaceaeD_5_Prevotellaceae UCG-001

D_0_BacteriaD_1_BacteroidetesD_2_BacteroidiaD_3_BacteroidalesD_4_RikenellaceaeD_5_Alistipes

D_0_BacteriaD_1_BacteroidetesD_2_BacteroidiaD_3_BacteroidalesOtherOther

D_0_BacteriaD_1_BacteroidetesD_2_CytophagiaD_3_CytophagalesD_4_CytophagaceaeD_5_Chryseolinea

D_0_BacteriaD_1_BacteroidetesD_2_CytophagiaD_3_CytophagalesD_4_CytophagaceaeD_5_Spirillum

D_0_BacteriaD_1_BacteroidetesD_2_FlavobacterialesD_3_FlavobacteriaceaeD_4_Flavobacteriaceae D_5_Flavobacterium

D_0_BacteriaD_1_ChlamydiaeD_2_ChlamydiaeD_3_ChlamydialesD_4_ParachlamydiaceaeD_5_Candidatus Prochlamydia

D_0_BacteriaD_1_ChloroflexiD_2_AnseroniaceaeD_3_AnseroniaceaeD_4_AnseroniaceaeD_5_Omnitella

D_0_BacteriaD_1_ChloroflexiD_2_TK100_3_uncultured bacteriumD_4_D_5_

D_0_BacteriaD_1_CyanobacteriaD_2_ChloroplastAmbiguous taxaD_4_D_5_

D_0_BacteriaD_1_CyanobacteriaD_2_ChloroplastOtherOtherOther

D_0_BacteriaD_1_CyanobacteriaD_2_MelainabacteriaD_3_GastreaerophilalesAmbiguous taxaD_5_

D_0_BacteriaD_1_CyanobacteriaD_2_MelainabacteriaD_3_GastreaerophilalesD_4_uncultured bacteriumD_5_

D_0_BacteriaD_1_CyanobacteriaD_2_MelainabacteriaD_3_GastreaerophilalesOtherOther

D_0_BacteriaD_1_CyanobacteriaD_2_VerrucomicrobialesAmbiguous taxaD_5_

D_0_BacteriaD_1_DerrifibacteresD_2_DerrifibacteresD_3_DerrifibacterialesD_4_DerrifibacteriaceaeD_5_Mucispilum

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_BacillalesD_4_AlcalobacillaceaeD_5_Turnebacillus

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_BacillalesD_4_BacillaceaeD_5_Bacilus

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_BacillalesD_4_Family XIIID_5_Engisubacterium

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_BacillalesD_4_PaenibacillaceaeD_5_Paenibacillus

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_BacillalesD_4_PlanococcaceaeAmbiguous taxa

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_BacillalesD_4_PlanococcaceaeD_5_Bhangavasea

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_BacillalesD_4_PlanococcaceaeD_5_Dombacillus

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_BacillalesD_4_PlanococcaceaeD_5_Kurtzia

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_BacillalesD_4_PlanococcaceaeD_5_Lysinibacillus

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_BacillalesD_4_PlanococcaceaeD_5_Rummelbacillus

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_BacillalesD_4_PlanococcaceaeD_5_Soliribacillus

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_BacillalesD_4_PlanococcaceaeD_5_Sporosarcina

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_BacillalesD_4_PlanococcaceaeOther

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_BacillalesD_4_StaphylococcaceaeD_5_Staphylococcus

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_LactobacillalesD_4_AerococcaceaeD_5_Aerococcus

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_LactobacillalesD_4_CarnobacteriaceaeD_5_Granulicatella

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_LactobacillalesD_4_CarnobacteriaceaeD_5_Trichococcus

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_LactobacillalesD_4_EnterococcaceaeD_5_Enterococcus

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_LactobacillalesD_4_EnterococcaceaeOther

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_LactobacillalesD_4_LactobacillaceaeD_5_Lactobacillus

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_LactobacillalesD_4_LeuconostocaceaeD_5_Leuconostoc

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_LactobacillalesD_4_LeuconostocaceaeD_5>Weissella

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_LactobacillalesD_4_Ro-042D_5_uncultured bacterium

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_LactobacillalesD_4StreptococcaceaeD_5Streptococcus

D_0_BacteriaD_1_FirmicutesD_2_BacilliD_3_LactobacillalesOtherOther

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_ChristensenellaceaeD_5_Christensenellaceae R-7 group

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_ChristensenellaceaeD_5_uncultured

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_ChristensenellaceaeOther

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_Clostridiaceae 1D_5_Candidatus Arthrornitis

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_Clostridiaceae 1D_5_Clostridium sensu stricto 1

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_Clostridiaceae 1D_5_Clostridium sensu stricto 8

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_Clostridiaceae 1Other

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_Clostridiales vadinBB60 groupAmbiguous taxa

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_Clostridiales vadinBB60 groupD_5_uncultured bacterium

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_Clostridiales vadinBB60 groupOther

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_DeftivallaceaeD_5_Deftivallaceae UCG-011

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_EubacteriaceaeD_5_Anaerofustis

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_Family XIIIID_5_Anaerovax

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_Family XIIIID_5_Family XIII AD3011 group

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_Family XIIIID_5_Family XIII UCG-001

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_Family XIIIID_5[Eubacterium] brachy group

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_Family XIIIID_5[Eubacterium] nodatum group

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_Family XIIIID_5Other

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_GracilibacteriaceaeD_5_uncultured

D_0_BacteriaD_1_FirmicutesD_2_ClostridiaD_3_ClostridialesD_4_LachnospiraceaeD_5_Acetofactor

D_0_BacteriaD_1_FirmicutesD_2