StaphGr-
Enterococcus
Streptococcus_parasuis_GCF_018987575.1_WP_016807377.1_23S_rRNA__adenine_2503_C_8__methyltransferase_Cfr_D_[Streptococcus_parasuis]
Akuthalobacillus clausii
Cohnella_thailandensis_GCF_017874075.1_WP_185122056.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Cohnella_thailandensis]
Cohnella_thailandensis_GCF_014212135.1_WP_185122056.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Cohnella_thailandensis]
Seinonella_peptonophila_GCF_900129355.1_WP_073150070.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Seinonella_peptonophila]
Bacillus_paramycoides_GCF_007682005.1_WP_098304113.1_MULTISPECIES_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_cereus_group]
Bacillus_cereus_GCF_002560375.1_WP_098304113.1_MULTISPECIES_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_cereus_group]
Anseuninibacillus_migulanus_GCF_000369965.1_WP_043068366.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Anseuninibacillus_migulanus]
Anseuninibacillus_migulanus_GCF_000369965.1_WP_043068366.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Anseuninibacillus_migulanus]
Anseuninibacillus_migulanus_GCF_001294525.1_WP_043068366.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Anseuninibacillus_migulanus]
Anseuninibacillus_migulanus_GCF_001294525.1_WP_043068366.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Anseuninibacillus_migulanus]
Anseuninibacillus_migulanus_GCF_001274715.1_WP_043068366.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Anseuninibacillus_migulanus]
Anseuninibacillus_migulanus_GCF_001274715.1_WP_043068366.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Anseuninibacillus_migulanus]
Sporosarcina_ginsengisoli_GCF_000427095.1_WP_025359241.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Sporosarcina_ginsengisoli]
Sporosarcina_psychrophila_DSM_447_GCF_000497245.1_WP_023511669.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Sporosarcina_psychrophila]
Cytobacillus_oceanisediminis_GCF_000704205.1_WP_053474707.1_MULTISPECIES_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillaceae]
Cytobacillus_solani_GCF_001420995.1_WP_053474707.1_MULTISPECIES_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillaceae]
Cytobacillus_sp._FJAT-21945_GCF_001275655.1_WP_053474707.1_MULTISPECIES_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillaceae]
Bacillus_gottheilii_GCF_000378125.1_WP_017155194.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus]
Bacillus_gottheilii_GCF_001499995.1_WP_017155194.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus]
Brevibacillus sp.
Brevibacillus_gravidiae
Sporosarcina_sp._Te-1_GCF_017499055.1_WP_207963382.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Sporosarcina_sp._Te-1]
Bacillus_sp._FJAT-49732_GCF_018343699.1_WP_213108897.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_sp._FJAT-49732]
Lysinibacillus_sphaericus_GCF_000378125.1_WP_013285616.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Lysinibacillus_sp._SOR037]
Bacillus_oleronius_GCF_017600215.1_WP_078109216.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_oleronius]
Bacillus_oleronius_GCF_002060555.1_WP_078109216.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_oleronius]
Bacillus_obstructivus_GCF_001897185.1_WP_071975457.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_obstructivus]
Bacillus_oleronius_GCF_018916995.1_WP_218391931.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_oleronius]
Bacillus_oleronius_GCF_018916995.1_WP_218391931.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_oleronius]
Bacillus_ginsengihumi_GCF_000427095.1_WP_088938831.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_ginsengihumi]
Paenibacillus_sp._FSL_R5-0345_GCF_000991425.1_WP_126999431.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Paenibacillus_sp._FSL_R5-0345]
Paenibacillus_polyxyma_GCF_000910975.1_WP_139025852.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Paenibacillus_sp._YH-14E5]
Paenibacillus_sp._J237S9_GCF_018403225.1_WP_213649291.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Paenibacillus_sp._J237S9]
Paenibacillus_sp._HJL_G12_GCF_000970355.1_WP_160490361.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Paenibacillus_sp._HJL_G12]
Bacillus_terrae_GCF_018333115.1_WP_212951417.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_terrae]
Bacillus_terrae_GCF_018333115.1_WP_212951417.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_terrae]
Bacillus_terrae_GCF_003605435.1_WP_120118601.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_terrae]
Lysinibacillus_sp._ZYM-1_GCF_001308875.1_WP_054809791.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Lysinibacillus_sp._ZYM-1]
Bacillus_nidoricus_GCF_000612855.1_WP_043476452.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_nidoricus]
Lysinibacillus_fusiformis_GCF_001726065.1_WP_069485093.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Lysinibacillus_fusiformis]
C dif&Enterococcus
Bacillus_sp._03113_GCF_000546885.1_WP_014430875.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_sp._03113]
Lysinibacillus_sp._FJAT-51151_GCF_016638705.1_WP_053394538.1_MULTISPECIES_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Lysinibacillus]
Lysinibacillus_sp._FJAT-14222_GCF_001278875.1_WP_053394538.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Lysinibacillus_sp._FJAT-14222]
Lysinibacillus_xylanilyticus_GCF_003903495.1_WP_110543767.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Lysinibacillus_xylanilyticus]
Bacillus pseudomycoides
Neobacillus_sp._ISL-48_GCF_018612965.1_WP_214601496.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_sp._ISL-48]
Bacillus_sp._ISL-51_GCF_018612635.1_WP_214601496.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_sp._ISL-51]
Bacillus_sp._ISL-26_GCF_018613395.1_WP_214601496.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_sp._ISL-26]
Bacillus_amyloliquefaciens_XHT_GCF_000221649.1_WP_013351126.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_amyloliquefaciens]
Bacillus_amyloliquefaciens_TA208_GCF_000190615.1_WP_013351126.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_amyloliquefaciens]
Bacillus_amyloliquefaciens_LL3_GCF_000190615.1_WP_013351126.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_amyloliquefaciens]
Bacillus_amyloliquefaciens_DSM_7_=_ATCC_23350_GCF_000190735.1_WP_013351126.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_amyloliquefaciens]
Bacillus_subtilis_GCF_000775125.1_WP_013351126.1_MULTISPECIES_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_subtilis_group]
Bacillus_amyloliquefaciens_GCF_013328839.1_WP_013351126.1_Cfr_family_23S_rRNA__adenine_2503_C_8__methyltransferase_[Bacillus_amyloliquefaciens]
Bacillus_amyloliquefaciens_GCF_000817555