

<p>ABC_tran GNR:HTH K09745.1:1:727954-732691 Bacillus cereus B4G1X1-1</p> <p>ERI8997.1 KE993461.1:48282-53536 Clostridiales bacterium oral taxon 876 str. F6540</p> <p>GBIC0680.1 BIST01000803.1:20473-26974 bacterium HRI7 Bacteria.</p> <p>HEY4680.1 DU5001000110.1:106-3699 Bacillota bacterium</p> <p>KAB814006.1 JXCB02000839.1:163876-170430 Tolypastris campylomonoides VBS12208</p> <p>KIM2708.1 J2UB1000114.1:16692-10973 Sulfolobum sp. FS86-10</p> <p>KKT79047.1 LX0201000825.1:16632-22559 Armatimonadetes bacterium CSPI-3</p> <p>MB2746078.1 JACLY010000533.1:1-4889 Flavobacterium sp.</p> <p>MBD1207289.1 JACZY010000011.1:51890-58284 Ignaviabacteria bacterium</p> <p>MBD325900.1 W1P01000322.1:16675-23787 Calditrixia sp.</p> <p>MBD300702.1 W1P01000160.1:2068-6943 Ignaviabacteria bacterium</p> <p>MBE047452.1 MJC0001000029.1:267-7824 Actinomycetota bacterium</p> <p>MBE218545.1 JACS5A01000038.1:1-5899 Rhodocyclales bacterium</p> <p>MBE057090.1 MJC0001000029.1:16286-40724 Actinomycetidium sp.</p> <p>MBE060205.1 5YSH01000006.1:24464-40343 Oscillospiraceae bacterium</p> <p>MBE012086.1 SV0301000011.1:21469-29503 Oscillospiraceae bacterium</p> <p>MBR002080.1 HEY01000002.1:226416-229907 Pseudomonas lundensis</p> <p>MBF080017.1 JAB210010000012.1:13066-10394 Clostridiales bacterium</p> <p>MB1307519.1 W1LV01000008.1:14051-12826 bacterium Bacteria.</p> <p>MBG38070.1 JACJ2010000120.1:138517-26254 Verrucomicrobia bacterium</p> <p>MBG58020.1 JACRLC010000003.1:10779-14175 Chloroflexi bacterium</p> <p>MBG02620.1 JACRLC010000003.1:1368-6640 Planctomycetota bacterium</p> <p>MBG07002.1 JACJ2010000008.1:105010-108577 Ignaviabacteria bacterium</p> <p>MBK710023.1 JAD30E01000008.1:109528-195404 Bacteroidetes bacterium</p> <p>MBK06031.1 JAD30E01000039.1:79446-84291 Rubrivivax sp.</p> <p>MBL048076.1 JAE552010000604.1:2772-8374 Nannocystaceae bacterium</p> <p>MBL06050.1 JAEURL010000605.1:718-5385 Amaelolineales bacterium</p> <p>MBL80754.1 JAEU0010000213.1:1-2575 Acidobacteria bacterium</p> <p>MBL02505.1 JAEUK010000063.1:115280-19092 Deintraprotobacteria bacterium</p> <p>MBL809412.1 JAEU3010000016.1:114-4920 Kinosporaceae bacterium</p> <p>MBN10202.1 JACG0100000158.1:1-6138 Thermoleptophila bacterium</p> <p>MBQ470543.1 JAF2UC010000101.1:1648-4453 Bacteroidales bacterium</p> <p>MBP010275.1 JAGNY010000003.1:13242-19088 Gammaproteobacteria bacterium</p> <p>MBQ24500.1 JAFPM010000155.1:1-3283 Acidaminococcaceae bacterium</p> <p>MBQ37532.1 JAFPM010000016.1:1156995-163482 Bacteroidales bacterium</p> <p>MBQ70008.1 JAFPM010000047.1:1306406-114786 Bacteroidales bacterium</p> <p>MBQ007005.1 JAFY0B010000052.1:11945-12364 Clostridia bacterium</p> <p>MBQ00046.1 JAFUR010000041.1:14013-8163 Phascolarctobacterium sp.</p> <p>MBQ08728.1 JAFM0100000514.1:1-3600 Oscillibacter sp.</p> <p>MBR17046.1 JAFVBB010000018.1:153263-162221 Bacteroidales bacterium</p> <p>MBR18106.1 JAFVBB010000003.1:18447-28841 Clostridia bacterium</p> <p>MBR52333.1 JAFY0B010000027.1:40802-49921 Clostridia bacterium</p> <p>MBR547243.1 JAFY0B010000093.1:218-5253 Oscillibacter sp.</p> <p>MBSH1280.1 JAFDV010000125.1:138564-23752 Acidobacteria bacterium</p> <p>MBSS0002.1 JACS2010000001.1:18672-26069 Actinomycetota bacterium</p> <p>MBUL17564.1 JAGW0Y010000061.1:6182-13757 Verrucomicrobia bacterium</p> <p>MBK03020.1 JANQV010000075.1:12339-15314 Gammaproteobacteria bacterium</p> <p>MEY070864.1 JALEPM010000018.1:164793-171367 Chitinophagaceae bacterium</p> <p>MBZ460217.1 JAIHAV010000060.1:1-6381 Thermotoga sp.</p> <p>MCAB0184.1 JACQ0010001538.1:1-1000 Candidatus Omnitrophota bacterium</p> <p>MCAN00008.1 JAGQ0B010000793.1:6714-12858 Mycococcales bacterium</p> <p>MCB010202.1 JAGQYF010000347.1:1-2129 Bacteroidetes bacterium</p> <p>MCQ77780.1 JAJDM010000002.1:1219-5486 Clostridiales bacterium</p> <p>MCJ270047.1 JAJTEZ010000119.1:1-3229 Saprospiraceae bacterium</p> <p>MCG315139.1 JAJVIR010000027.1:78875-87492 bacterium Bacteria.</p> <p>MC418088.1 JACW0Y010000045.1:1-4232 Acidaminococcaceae bacterium</p> <p>MCJ00009.1 JALFCR010000171.1:1-2689 Clostridiales bacterium</p> <p>MCL20600.1 W1RL010001041.1:13988-29704 Oscillospiraceae bacterium</p> <p>MCMA00418.1 JANQD010000073.1:3135-7250 Actinoplanes hotanensis</p> <p>MCO2000413.1 JACS0M010000012.1:146359-56425 Bacteroidales bacterium</p> <p>MCQ259049.1 JACS0M010000132.1:1152-4301 Phascolarctobacterium sp.</p> <p>MCS0000.1 V1L010000004.1:140247-46555 Acidaminococcus fermentans</p> <p>MST94547.1 S1B01000038.1:202-6388 Pedosphaera sp.</p> <p>MYR4005.1 VYANO10000412.1:1-3353 Candidatus Poribacteria bacterium</p> <p>NCC37726.1 RZQ010010739.1:133-1012 Chloroflexia bacterium</p> <p>NCO60003.1 JACCTV010000008.1:10324-18588 Candidatus Aquasulter secundus</p> <p>NAG0000.1 W00V01000136.1:9074-16795 Armatimonadota bacterium</p> <p>NMA043773.1 JABEWH010000158.1:14092-7668 Chlamydiae bacterium</p> <p>NQJ02184.1 JABP01010000001.1:30613-48015 Coralibacillus coralitoides</p> <p>NPV0046.1 JANL0W010000020.1:147852-151397 Bacillota bacterium</p> <p>OCB67429.1 M6S010000044.1:15583-8920 Calditrixia sp. RW_13_4_0</p> <p>PCJ20376.1 WY0V010001000.1:148454-144906 Candidatus (o)acmonadota bacterium</p> <p>QYQ000.1 C972784.1:1246174-125265 Aminithiophilus ramosus</p> <p>RMB11096.1 REFR010000089.1:325583-334450 Eliatimonas nillapovae</p> <p>RMC02620.1 RHM010000092.1:1-2248 Armatimonadetes bacterium</p> <p>TCQ27004.1 SHZE01000053.1:25-2190 Hydromonas duriensis</p> <p>TMPP7177.1 V00001000352.1:3589-6481 Verrucomicrobia bacterium</p> <p>WP_111910208.1 NZ_JACTY0401000024.1:1:43-2262 Acinetobacter baumannii</p> <p>WP_01530862.1 NC_020126.1:15</p>
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