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Bacillus cereus E331

AGE03047.1  
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Bacillus subtilis XF-1

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CP067699.2:60826-69082  
Streptomyces lydicus

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Mycobacterium

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AN000010000008.1:214239-215498  
Salmonella enterica

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AAPGV010000213.1:1-1045  
Salmonella enterica subsp. enterica serovar Kentucky

EHK80898.1  
JAK001000129.1:106924-22114  
Saccharomonospora azurea Z21C  
14680

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JEW01000032.1:36674-39812  
Acinetobacter baumannii 295505

GHP10057.1  
BNB01000028.1:21414-26114  
Streptomyces thermotolasticus

HHH28774.1  
DRO101000562.1:275-2967  
Polysphingaceae bacterium

KAA0215069.1  
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Cyanobacteria bacterium CY4

KAR057807.1  
VZP081000086.1:1-2116  
Ideonella dechloratans

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candidate division TM6 bacterium GM2011\_GMF2\_36\_131

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JABUR01000046.1:565-16513  
Mycobacter sp.

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Deltaproteobacteria bacterium

MBK608101.1  
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MBK815291.1  
JAD30061000006.1:383425-386263  
Saprospiraceae bacterium

MBK826685.1  
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Polysphingaceae bacterium

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MBK9453870.1  
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Bacteroidetes bacterium

MBQ4031662.1  
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Bacilli bacterium

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Lachnospiraceae bacterium

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Nitrosomonas europaea

MBX306660.1  
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Streptomyces sp. MNC04124

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Planctomycetaceae bacterium

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Bacteroidia bacterium

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Epsilonproteobacteria bacterium

MC0101069.1  
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Paraburkholderia sediminicola

MC08031280.1  
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Lentisphaeraceae bacterium

MDH643145.1  
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Cronobacter dublinensis

MSR27083.1  
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Candidatus Armanicantae bacterium

NJM23481.1  
JAIU1010000479.1:12233-5079  
Rickettsia sp. SM\_7\_0

OBX76551.1  
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Moraxella atlantae

QJW4709.1  
MOL010000067.1:15979-33519  
Alphaproteobacteria bacterium 41\_20

OC044703.1  
MLH031000045.1:3218-7697  
Rodentibacter trehalosifermentans

OQA30359.1  
PMB021000029.1:77-8564  
Candidatus Dependulae bacterium A0urb.B10331

OZC17032.1  
LSA031000074.1:103626-106902  
Sporomusa silvatica DSM 10669

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Flavobacterium columnare NBRC 100251 = ATCC 23463

PJK1125.1  
NMR021000081.1:402477-410050  
Lyobacteraceae bacterium MML08\_0793

QIO1075.1  
CP049316.1:1677827-1680612  
Acinetobacter lantii

QMU5413.1  
CP046177.1:2974552-2977351  
Flavobacteriaceae bacterium

RQR0253.1  
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Burkholderia sp. B0902

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Chitinophagaceae bacterium

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Flavobacterium sp.

RZQ0065.1  
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Anoxytosis Suedae

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Streptomyces sp. DconL5

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Paracitrovirax valerianellae

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Fontibacillus panacineptis

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Streptomyces zhaozhouensis

TAG05096.1  
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Oscillatoriales cyanobacterium

TWK74524.1  
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Bacillus licheniformis

TWK80549.1  
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Nitrilobacillus boronitolerans F182

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Proteobacteria sp. oral taxon 317 str. F0108

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Carnocytophaga sp. oral taxon 332 str. F0301

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Flavobacterium branchiophilum FL-15

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Bacillus velezensis

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Gemmatirosa kalamazooensis

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NZ\_KK211315.1:1607-4897  
Butyrivibrio sp. MC2021

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NZ\_J0001000028.1:8516-20754  
Halaea salicispens DSM 19537

WP\_05216434.1  
NZ\_JMK01000021.1:102239-109029  
Lysobacter dokdonensis 05-58

WP\_150000041.1  
NZ\_LD0031000011.1:1-388  
Streptomyces sp. W0054

WP\_234305127.1  
NZ\_P0V01000008.1:75673-78990  
Meisseria sicca

WP\_180761017.1  
NZ\_JADHT010000009.1:83977-98406  
Pseudomonas palleroniana

WP\_176702113.1  
NZ\_L76702100034.1:129758-132393  
Gilliamella sp. Fert-1

WP\_02706128.1  
NZ\_J0001000028.1:8516-20754  
Halaea salicispens DSM 19537

WP\_05216434.1  
NZ\_JMK01000021.1:102239-109029  
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