

AY125958.1:1:1082954-1086498  
*Pseudomonas aeruginosa*

CCG18016.1  
HEB41473.1:1783002-787508  
*Taylorella equigenitalis* 14/56

EEY51416.1  
GC104315.1:1:214561-217912  
*Pasteurella dagmatis* ATCC 43325

EJ105708.1  
CM011559.1:1:2737811-2742283  
*Pseudomonas chlororaphis* subsp. *aureofaciens* 30-84

ETD72835.1  
AYS01000015.1:152-3467  
*Pelistega indica*

EYA71780.1  
JGE001000038.1:18429-22832  
*Bacteroides fragilis* str. S24L15

HBK05410.1  
DCC001000151.1:12113-8496  
*Bacteroides bacterium*

HDQ46180.1  
DSAB1473.1:1:4756-7857  
*bacterium* *Bacteria*.

HJC17833.1  
DWF010000271.1:97324-102467  
*Candidatus Allostipes stercorigallinarum*

KAF0203985.1  
WSK01000004.1:1:98787-107857  
*Bacteroidetes bacterium*

KHF30061.1  
JRLC01000006.1:1:39690-42656  
*Anoxybacillus* sp. *BC01*

KKB45390.1  
KQB33924.1:1:1-1565  
*Parabacteroides gordonii* MS-1 = *DSM 23371*

MBE485008.1  
JACHJ010000033.1:1:204-9101  
*Streptomyces griseomycini*

MBC24118.1  
PAB101000029.1:1:24679-29957  
*Phycisphaerae bacterium*

MBD6708743.1  
JACY1010000018.1:1:154032-162657  
*Pseudomonas* sp. *CFBP 13711*

MBF1065876.1  
JABZ0010000009.1:1:724-4545  
*Prevotellaceae bacterium*

MBI1934319.1  
JACU0010000075.1:1:94-2801  
*Candidatus Peregrinibacteria bacterium*

MB1235690.1  
JACPX010000029.1:1:42866-47895  
*Candidatus Peregrinibacteria bacterium*

MB1685357.1  
JAC02001000077.1:1:1972-8909  
*Candidatus Wallibacteria bacterium*

MB156188.1  
JAC10010000090.1:1:1284-5361  
*Candidatus Peregrinibacteria bacterium*

MBK8402491.1  
JAD7AT010000015.1:1:776531-783639  
*Protonivibrio* sp.

MBN2712280.1  
JAFGW0100000215.1:1:14902  
*Planctomycetota bacterium*

MBN8703277.1  
JAFLBG010000040.1:1:5451-9523  
*Bacteroidetes bacterium*

MBQ090303.1  
JAFPM010000258.1:1:36-3689  
*Solirubrobacterales bacterium*

MBQ7166493.1  
JAC02001000052.1:1:2004-6670  
*Kiritimatiellae bacterium*

MBQ7628725.1  
JAC10010000124.1:1:241-7655  
*Bacteroidales bacterium*

MBP550538.1  
JAGCV0100000012.1:1:1-10096  
*Bacteroidales bacterium*

MBP566346.1  
JAGCS0100000065.1:1:147-7171  
*Bacteroidetes bacterium*

MBP690464.1  
JAG0PE010000006.1:1:6862-14283  
*Candidatus Paceibacterota bacterium*

MBQ624246.1  
JAFVW010000057.1:1:12379-15203  
*Bacteroidaceae bacterium*

MBQ711745.1  
JAFUT010000063.1:1:1-8516  
*Prevotella* sp.

MBQ954824.1  
JAFTE010000020.1:1:10663-30568  
*Bacteroidales bacterium*

MBR346091.1  
JAFWQ0100000026.1:1:69378-79054  
*Lachnospiraceae bacterium*

MBR3896723.1  
JAFWQ0100000076.1:1:18567-25261  
*Bacteroidaceae bacterium*

MBR3067157.1  
JAFXBA010000054.1:1:30629-33109  
*Bacteroidales bacterium*

MBT3202020.1  
JABG00100000001.1:1:37471-40839  
*Candidatus Peregrinibacteria bacterium*

MBV9320009.1  
JAF4V010000051.1:1:14535-19019  
*Acidobacteriaceae bacterium*

MBX321740.1  
JAFBMA010000167.1:1:1-6842  
*Labililthrix* sp.

MBX3765185.1  
QGN01000238.1:1:62-683  
*Burkholderia cepacia*

MCA6492120.1  
JADBXE010000057.1:1:159-3877  
*Chitinophagaceae bacterium*

MCA915730.1  
JAG0P0100000070.1:1:762-7557  
*Planctomycetales bacterium*

MCA070563.1  
JAG011010001376.1:1:9780-19053  
*Mycococcales bacterium*

MCB1600976.1  
JAF031010000171.1:1:1477-5799  
*Lysobacterales bacterium*

MCB9382367.1  
JACJWP0100000015.1:1:368-8464  
*Acidimicrobiaceae bacterium*

MCC6741780.1  
JADLHC0100000307.1:1:1989-3662  
*Planctomycetota bacterium*

MCD9583380.1  
JAJQXT010000105.1:1:1-1352  
*Tenacibaculum maritimum*

MDF0607206.1  
JACCFB020000062.1:1:736-3459  
*Neisseriaceae bacterium* *TCSR-5*

MC03002.1  
RZZ031000494.1:1:1-2025  
*Opitutia bacterium*

NEZ59010.1  
QW001000004.1:1:120706-123137  
*Adonisia turfae* *CM09081*

NJK31821.1  
JAAU0010000140.1:1:381-4942  
*Beltaproteobacteria bacterium*

NLE21221.1  
JAAZAR0100000006.1:1:679-6697  
*Actinomycetota bacterium*

NUO72258.1  
JABFX010000192.1:1:5768-8298  
*Fraternia* sp.

NYH51431.1  
JACHL0100000001.1:1:1181975-1185409  
*Nocardopsis sinuspersici*

OGU73714.1  
HMA101000036.1:1:1-5856  
*Ignavibacteria bacterium* *RIFCSFLOW2\_02\_FULL\_55\_14*

OIO54398.1  
HMM001000020.1:1:134488-138427  
*Candidatus Peregrinibacteria bacterium* *CG1\_02\_54\_53*

ORF02093.1  
MAG010000052.1:1:2536-6198  
*Snodgrassella alvi*

PMY32656.1  
POGF01000046.1:1:27140-31718  
*Pseudomonas* sp. *GW56-L14*

PTV55566.1  
QAY001000137.1:1:372-3829  
*Pseudomonas putida*

QEV73709.1  
CP023688.1:1:18058-26089  
*Streptomyces rimosus*

QGS44403.1  
CP045378.1:1:1450823-1454647  
*Streptococcus gordonii*

QOW62145.1  
CP051839.1:1:1160841-1164311  
*Treponema pedis*

RME20003.1  
RFKJ01000035.1:1:1-3737  
*Beltaproteobacteria bacterium*

RUM53976.1  
QOAT01000172.1:1:2736-5695  
*Thiothrix* sp.

SUB94547.1  
UGTN010000002.1:1:961979-971134  
*Prevotella denticola*

VUD51028.1  
CABFF0100000003.1:1:9438-18214  
*Thalassocella blandensis*

WP\_010539506.1  
NZ\_LJG001000051.1:1:5403  
*Bacteroides faecis* *MAJ27*

WP\_123431118.1  
NZ\_JAMKE010000004.1:1:534255-542031  
*Pseudomonas* sp. *BLG00176*

WP\_098793255.1  
NZ\_NUNC01000038.1:1:128444-124112  
*Bacillus cereus*

WP\_167375423.1  
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*Pseudomonas agarrici*

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*Photorhabdus lumumidii* subsp. *clarkiei*

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*Actinolioteichus* sp. *GBA129-24*

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*Xylaniobacter brevis* *P6B11*

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*Pseudomonas syringae* pv. *atrofaciens*

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