DQ856526.1 gg_id:246512 intestinal microflora Chinese mitten crab (Eriocheir sinensis) clone C2F DQ856
EF157278.1 gg_id:258039 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF1572
U90216.1 gg_id:6294 Sphingomonas sp. 2MPII U9021
EU682685.1 gg_id:342749 Sphingomonas sp. str. V2M44 EU682
GQ020378.1 gg_id:489291 Topographical and Temporal Human Skin Microbiome skin glabella clone nbu268d11 GQ020
U20775.2 gg_id:44107 Novosphingobium stygium str. SMCC B0712 U2077
EU335142.1 gg_id:257791 Changes microbial metabolic and along hydrogeochemically variable profile saturate EU335
DQ906835.1 gg_id:256780 Oman subsurface soil clone 7B-8 DQ906
GU166344.1 gg_id:582973 Altererythrobacter sp. str. JM27 GU166
EU707558.1 gg_id:352352 Novosphingobium sp. str. YC6720 EU707
AJ746092.1 gg_id:113876 Novosphingobium sp. MG35 AJ746
EU726272.1 gg_id:349727 Alterierythrobacter sp. str. H32 EU726
FJ542890.1 gg_id:525932 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone A FJ5428
DQ497759.1 gg_id:161477 Sphingomonas sp. str. M80C3 DQ497
DQ295907.1 gg_id:147648 Cohn's Crenothrix filamentous oxidizer unusual methane monooxygenase physical e DQ295
GU180153.1 gg_id:555283 Impact coplanar PCBs on microbial activity anaerobic estuarine sediments sediment GU180
Y13774.1 gg_id:6269 Blastomonas natatoria str. DSM 3183 Y13774
AM940945.1 gg_id:282135 Sphingomonas sp. ECN-2008 AM940
D13737.1 gg_id:6255 Sphingomonas suberifaciens str. IFO 15211 D13737
EU202871.1 gg_id:340768 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU2028
EF516033.1 gg_id:222080 grassland soil clone FCPS756 EF5160
EF580938.1 gg_id:278358 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM4 EF5809
GQ008488.1 gg_id:501177 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ008
FJ478827.1 gg_id:591215 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4788
EU202876.1 gg_id:304392 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU2028
DQ248234.1 gg_id:143386 carbon tetrachloride contaminated soil clone UE1 DQ248
EU223953.1 gg_id:249385 -RFLP microbes Qinghai-Tin rail soil clone 120 EU2239
EU193103.1 gg_id:367223 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU193
AY922159.1 gg_id:114292 farm soil clone AKYH831 AY922
EU193101.1 gg_id:367335 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU193
AY850294.1 gg_id:136384 depending on what microbial treatment present soil clone w4 AY8502
EU193107.1 gg_id:364127 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU193
AB258386.1 gg_id:159768 Kaistobacter terrae str. KCTC12630 AB2583
EU202894.1 gg_id:344045 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU2028
AY769084.1 gg_id:108378 Kaistobacter koreensis str. PB180 AY7690
EU202882.1 gg_id:340856 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU2028
AM935913.1 gg_id:563989 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM935
EU133392.1 gg_id:242675 tallgrass prarie soil clone FFCH13278 EU133
GQ051344.1 gg_id:414905 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu4 GQ051
FJ266331.1 gg_id:549411 Novosphingobium capsulatum str. RFNB21 FJ2663
AB196250.1 gg_id:115182 Sphingomonas sp. str. IC081 AB196

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AB275605.1 gg_id:214353 Sphingomonas bosoensis str. WF3 AB275
AB360760.1 gg_id:240554 Novosphingobium sp. str. NIY3 AB360
AB088760.1 gg_id:78821 Novosphingobium sp. str. J30 AB088
EU304250.1 gg_id:248028 Novosphingobium str. Hg4-04 EU304
AB453304.1 gg_id:514645 Sphingomonas sp. str. SO11 AB453
GQ388869.1 gg_id:511687 quality deterioration drinking system during serious red water outbreak clone Z28 GQ388
EU730906.1 gg_id:332390 Novosphingobium stygium str. 159 EU730
DQ137853.2 gg_id:138108 Novosphingobium sp. str. Geo25 DQ137
GQ388914.1 gg_id:580916 quality deterioration drinking system during serious red water outbreak clone Z76 GQ388
FJ425737.1 gg_id:364924 Novosphingobium sp. str. CC-TPE-1 FJ4257
DQ337059.1 gg_id:149880 subsurface water clone EV818SWSAP5 DQ337
GQ264150.1 gg_id:579694 Structure Simulated Level Waste Site simulated level waste site clone WC3_91 GQ264
GQ023414.1 gg_id:486255 Topographical and Temporal Human Skin Microbiome skin buttock clone nbu318b1 GQ023
U20755.2 gg_id:73612 Novosphingobium aromaticivorans str. SMCC B0695 U20755
AB362778.1 gg_id:256401 Sphingomonas sp. str. YT0136 AB362
GQ103626.1 gg_id:438411 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ103
GQ069267.1 gg_id:374280 Topographical and Temporal Human Skin Microbiome skin buttock clone nbw241e0 GQ069
AY571829.1 gg_id:102884 Sphingomonas sp. 44/40 AY5718
AY792290.1 gg_id:127870 Humic Lake clone CrystalBog2A11 AY792
FJ377543.1 gg_id:369876 Novosphingobium sp. str. F2 FJ3775
GQ038636.1 gg_id:412772 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu4 GQ038
GQ020999.1 gg_id:488670 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu2 GQ020
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GQ021683.1 gg_id:487986 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ021
GQ472779.1 gg_id:576436 Biodiversity planktonic Northern surface water northern Bering Sea clone u3 GQ472
FJ459983.1 gg_id:453127 Novosphingobium subterraneum str. C1-3-2; IMER-A3-1 FJ4599
GQ038752.1 gg_id:378163 Topographical and Temporal Human Skin Microbiome skin buttock clone nbu522f03 GQ038
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EU937862.1 gg_id:313934 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU937
AY921998.1 gg_id:113391 farm soil clone AKYG1872 AY921
EU644200.1 gg_id:281742 and Samoylov Island Lena active layer -centred polygon near surface horizon; Rim_T EU644
AJ416411.1 gg_id:38548 Novosphingobium hassiacum W-51 AJ4164
AB177224.1 gg_id:133463 seafloor methane hydrate clone ODP1230B9.15 AB177
EF127604.1 gg_id:208477 Fossil Genes and Microbes Entombed Oldest Ice on Earth ice clone EME109 EF1276
GQ083294.1 gg_id:403709 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ083
AF411072.1 gg_id:39863 Sphingomonas sp. str. ACM-3962 AF411
GQ038964.1 gg_id:371390 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu4 GQ038
GQ102705.1 gg_id:382549 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ102

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DQ327674.1 gg_id:165838 Biological filtration limits carbon availability affects downstream biofilm formation a DQ327
GQ023478.1 gg_id:486191 Topographical and Temporal Human Skin Microbiome skin buttock clone nbu319b0(GQ023
GQ092194.1 gg_id:449436 Topographical and Temporal Human Skin Microbiome skin palm clone nbw409f03c1 GQ092
EF549586.1 gg_id:252275 Novosphingobium indicum str. H25 EF5495
AY328849.1 gg_id:90554 drinking water MB27 AY3288
FJ801185.1 gg_id:513803 Investigation Water -DGGE and Clone Library Zoige Alpine Wetland clone ZWB2-14 FJ8011
GQ099344.1 gg_id:439803 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw5(GQ099
EU431800.1 gg_id:562657 Troglobitic (Cansiliella) on specialized feeding apparatus calcium carbonate (moonmi EU4318
AB240479.1 gg_id:142028 microbial structure rhizosphere biofilm Sapporo root-tip (0 40 mm) Phragmites Sosei AB2404
DQ463204.1 gg_id:157220 Acid Drainage Tong Lushan Copper Mine China acid mine drainage clone tls3-9 DQ463
GQ022145.1 gg_id:487522 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu3 GQ022
AF025352.1 gg_id:6293 Sphingomonas sp. str. D16 AF0253
EF029110.1 gg_id:221739 Novosphingobium resinovorum str. NCIMB 8767 EF0291
DQ450761.1 gg_id:160417 Microbial Alpine Tundra Wet Meadow Soil: Chloroflexi Cold Water-Saturated saturat DQ450
EU639137.1 gg_id:323096 Ecological Role Firmicutes Identified Thermophilic Microbial Fuel Cells thermophilic n EU6391
DQ501315.1 gg_id:189621 Seasonal dynamics free-living and particle- four lakes northeastern Germany freshw DQ501
FJ801188.1 gg_id:542132 Investigation Water -DGGE and Clone Library Zoige Alpine Wetland clone ZWB3-1 FJ8011
GQ020347.1 gg_id:489319 Topographical and Temporal Human Skin Microbiome skin glabella clone nbu267h11 GQ020
U20756.2 gg_id:136117 Novosphingobium aromaticivorans str. SMCC F199 U20756
DQ676432.1 gg_id:224427 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676
FM872861.1 gg_id:360253 occupant source house floor dust clone FB03H05 FM872
FM178802.1 gg_id:311289 Monitoring abundance and expression 'dehalococcoides' species chloroethene-redu FM178
AY792291.1 gg_id:150609 Temporal Variation and Humic Lake clone CrystalBog022E8 AY7922
DQ207361.1 gg_id:139541 Sphingomonas sp. str. PF-I DQ207
EU104173.1 gg_id:341018 Evidence Comamonadaceae determinants settling performance activated sludge clo EU1041
CU917745.1 gg_id:533625 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU9177
EU827494.1 gg_id:352634 Sphingomonas sp. tfd44 str. TFD44 EU8274
AF097802.2 gg_id:21583 activated sludge clone 1958 AF0978
AY212578.1 gg_id:99539 water 10 m downstream manure clone 127ds10 AY2125
GQ050992.1 gg_id:435628 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ050
AB286635.1 gg_id:214915 Reducing production and domination comamonadaceae reducing oxygen supply was AB2866
AY922004.1 gg_id:113584 farm soil clone AKYG395 AY9220
FJ177534.2 gg_id:341537 Novosphingobium sp. str. HU1-AH51 FJ1775
GQ069243.1 gg_id:397437 Topographical and Temporal Human Skin Microbiome skin buttock clone nbw241b0 GQ069
GQ037883.1 gg_id:464605 Topographical and Temporal Human Skin Microbiome skin interdigital web space clc GQ037
AJ292613.1 gg_id:150636 Combined use and study polychlorinated biphenyl-polluted soil clone WD2119 AJ2926
Z93995.1 gg_id:6292 activated sludge clone T28 Z93995
U20774.2 gg_id:70401 Novosphingobium aromaticivorans str. SMCC B0522 U20774
AB237654.1 gg_id:156847 Novosphingobium subarcticum str. T7b AB2376
AB177883.1 gg_id:103401 Novosphingobium sp. str. TUT562 AB1778

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GQ093042.1 gg_id:426566 Topographical and Temporal Human Skin Microbiome skin elbow clone nbw408d11 GQ093
GQ022398.1 gg_id:487271 Topographical and Temporal Human Skin Microbiome skin axillary vault clone nbu31 GQ022
GU208446.1 gg_id:563756 Study on Sediment Dongping Lake sediment clone Fr2-8 GU208
GQ008813.1 gg_id:500852 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ008
AY328817.1 gg_id:90016 drinking water Y14 AY328
DQ337064.1 gg_id:146133 subsurface water clone EV818SWSAP16 DQ337
EU937930.1 gg_id:348236 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU937
AY500142.1 gg_id:104274 Novosphingobium sp. T3-B9 str. T3-B9; AS 1.3432 AY500
EU907884.1 gg_id:320881 Identification hexahydro-135-trinitro-135-triazine (RDX)-degrading microorganisms v EU907
EU431820.1 gg_id:554799 Troglobitic (Cansiliella) on specialized feeding apparatus calcium carbonate (moonmi EU431
EU984513.1 gg_id:306710 Novosphingobium sp. str. Rr 2-17 EU984
GQ035853.1 gg_id:418017 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbu37 GQ035
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AB074191.1 gg_id:24166 Sphingomonas sp. str. HL7 AB074
EU937960.1 gg_id:342517 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU937
GQ007148.1 gg_id:502513 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ007
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AB491315.1 gg_id:546882 Sphingobium fuliginis str. TIK-1 AB491
AB042233.1 gg_id:18499 Sphingobium herbicidovorans AB042
EF065102.1 gg_id:179619 Sphingobium herbicidovorans str. FL EF065
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DQ885954.1 gg_id:162032 Sphingomonas sp. str. BSL8 DQ885
AY574367.1 gg_id:106220 Sphingobium yanoikuyae str. DD109 AY574
AB331239.1 gg_id:240760 Sphingobium yanoikuyae str. FM-3 AB331
Y12803.1 gg_id:38158 Sphingomonas agrestis str. HV3 Y12803
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AB025530.1 gg_id:6164 Sphingomonas sp. str. MBIC3990 Lo251 AB025
AM489507.1 gg_id:203347 Sphingomonas olei str. IMMIB HF-1 AM489
GQ025136.1 gg_id:415594 Topographical and Temporal Human Skin Microbiome skin axillary vault clone nbw1 GQ025
EU744623.1 gg_id:349718 Sphingobium sp. str. W1 EU744
AF157490.1 gg_id:6196 Sphingomonas sp. str. JS5 AF157
EU050705.1 gg_id:238200 Microbial waters production water Daqing oil field clone DQ328-85 EU050
EU286535.1 gg_id:239979 Sphingobium sp. 2F5-2 EU286
AB112465.1 gg_id:88144 Suiyo Seamount hydrothermal plume clone SUP27-27 AB112
AY771799.1 gg_id:108045 Sphingomonas sp. 12-7 AY771
GQ096356.1 gg_id:423731 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw4 GQ096
DQ202286.1 gg_id:139757 Sphingomonas sp. str. PF-K DQ202
AB000106.1 gg_id:6183 Sphingomonas sp. str. VT1 AB000

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AB299582.1 gg_id:254239 Sphingomonas sp. str. M30-VN10-1W AB299
FJ192049.1 gg_id:315748 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1920
EU159273.1 gg_id:354715 Sphingobium sp. str. YBL3 EU159
D13728.1 gg_id:6148 Sphingobium yanoikuyae str. IFO15102 D13728
EU009209.1 gg_id:255180 Sphingobium yanoikuyae str. WP01 EU009
X94098.1 gg_id:6200 Sphingomonas xenophaga str. BN6 X94098
GQ086522.1 gg_id:467619 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw29 GQ086
AY367025.1 gg_id:93363 Sphingomonas sp. As1-3 AY367
X94100.1 gg_id:6163 Sphingomonas paucimobilis str. EPA 505 X94100
AF177917.1 gg_id:6193 Sphingomonas sp. str. DhA-95 AF177
AF191022.1 gg_id:6167 Sphingomonas sp. str. CD AF191
EU183233.1 gg_id:251876 Sphingobium yanoikuyae str. XJ EU183
EU307932.1 gg_id:239648 Sphingobium yanoikuyae str. BF-18 EU307
FJ191324.1 gg_id:310721 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1913
EU375008.1 gg_id:275989 Sphingomonas sp. str. PhS5B EU375
AM998824.1 gg_id:552210 Sphingomonas sp. VC-230 str. VC-230=CCUG 55809 AM998
X85023.1 gg_id:6161 Sphingobium yanoikuyae str. PAH-degrading X85023
AJ011505.1 gg_id:6162 Sphingomonas sp. str. DhA-33 AJ0115
FJ191470.1 gg_id:332159 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1914
AJ000918.2 gg_id:40332 Sphingomonas sp. str. K6 AJ0009
FJ191669.1 gg_id:329197 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1916
EU240401.1 gg_id:252134 Sphingomonas xenophaga str. B107 EU240
AY367026.1 gg_id:93744 Sphingomonas sp. As1-5 AY367
FJ966197.1 gg_id:589286 Sphingobium sp. str. P21 FJ9661
FJ193541.1 gg_id:322175 Genetic inventory 'LMA-MTF' spacecraft assembly clean room during Phoenix missior FJ1935
U37341.1 gg_id:6178 Sphingomonas paucimobilis U3734
AY635898.1 gg_id:101815 Sphingomonas sp. str. AMS7 AY6358
EU616621.1 gg_id:271841 Sphingomonas sp. str. JK1 soil EU616
U37525.1 gg_id:6160 Sphingobium yanoikuyae str. Q1 U37525
EU159274.1 gg_id:313442 Sphingobium sp. str. YBL1 EU159
D16145.1 gg_id:6171 Sphingobium yanoikuyae str. GIFU9882 D16145
GQ019477.1 gg_id:490192 Topographical and Temporal Human Skin Microbiome skin umbilicus clone nbu190d GQ019
GQ101175.1 gg_id:370761 Topographical and Temporal Human Skin Microbiome skin manubrium clone nbw49 GQ101
AY151393.1 gg_id:73215 Sphingomonas cloacae LH227 AY1513
AB299577.1 gg_id:251303 Sphingomonas sp. str. M5-VN5-4W AB299
AB214867.1 gg_id:131055 Sphingomonas sp. str. ED8 AB2148
FJ191966.1 gg_id:306789 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1919
GQ087461.1 gg_id:416479 Topographical and Temporal Human Skin Microbiome skin buttock clone nbw357g0 GQ087
GQ023300.1 gg_id:486369 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu2 GQ023

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EU057882.1 gg_id:253243 Sphingomonas sp. CH30-8 EU0578
AB040739.1 gg_id:41591 Sphingomonas cloacae str. S-3 AB0407
EU095328.1 gg_id:250814 Sphingobium sp. str. x23 EU0953
AJ009708.2 gg_id:51138 Sphingobium sp. K40 AJ0097
AB120764.1 gg_id:94883 Sphingobium yanoikuyae str. EC-S01 AB1207
EU219973.1 gg_id:251683 Sphingomonas sp. str. KMC001 EU2199
AF038030.1 gg_id:6195 Sphingomonas sp. str. Lep1 AF0380
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GQ036338.1 gg_id:409030 Topographical and Temporal Human Skin Microbiome skin palm clone nbu363b03c1 GQ036
AY921715.1 gg_id:113008 farm soil clone AKYH784 AY9217
EU240418.1 gg_id:257571 Sphingobium yanoikuyae str. B166 EU2404
AY151392.1 gg_id:55376 Sphingomonas chungbukensis VM0440 AY1513
FJ193550.1 gg_id:307264 Genetic inventory 'LMA-MTF' spacecraft assembly clean room during Phoenix missior FJ1935
AB214868.1 gg_id:131403 Sphingomonas sp. str. ED9 AB2148
AF494538.1 gg_id:67345 Sphingomonas sp. str. BPC4 AF4945
AY328789.1 gg_id:90697 drinking water system simulator clone DSSD92 AY3287
AY519129.1 gg_id:99094 Sphingomonas paucimobilis str. B90A AY5191
AB288318.1 gg_id:212603 Sphingomonas sp. str. SY-17 AB2883
AY921943.1 gg_id:113573 farm soil clone AKYG1014 AY9219
AY026948.1 gg_id:32571 Sphingomonas sp. str. ML1 AY0269
GQ087010.1 gg_id:432007 Topographical and Temporal Human Skin Microbiome skin axillary vault clone nbw3 GQ087
EU580697.1 gg_id:261294 Sphingobium sp. str. NP22b EU5806
DQ238831.1 gg_id:142891 Sphingobium sp. VM2 DQ238
AM935351.1 gg_id:534250 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM935
GQ022171.1 gg_id:487497 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu3 GQ022
FJ192016.1 gg_id:324267 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1920
FJ192036.1 gg_id:350560 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1920
AY646154.1 gg_id:136602 Sphingomonas sp. str. 3Y AY6461
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FJ796422.1 gg_id:462836 Sphingomonas sp. str. S-II-13 FJ7964
AJ007800.1 gg_id:6234 Asticcacaulis excentricus str. ACM 1263 (T) AJ0078
AF331660.1 gg_id:136175 Caulobacter leidyia AF3316
FN421869.1 gg_id:593077 proteogenomics physiology phyllosphere clover clone 2_E01 FN4218
AY571825.1 gg_id:102263 hydrocarbon-contaminated soil clone 433E AY5718
FJ605269.1 gg_id:560773 Sphingomonas sp. str. DCY44 FJ6052
AM229669.1 gg_id:152429 Sphingomonas mucosissima str. CP173-2 AM229
AF409011.1 gg_id:102915 Sphingomonas sp. str. Ellin169 AF4090
GU377113.1 gg_id:523159 Sphingomonas sp. str. sptzw24 GU377
FN421984.1 gg_id:585288 proteogenomics physiology phyllosphere clover clone 5_F05 FN4219

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EU528244.1 gg_id:285960 Seasonal Dynamics Mudflat Mouth Major Kentucky Lake Reservoir Tributary sedime EU528
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NR_028634.1 gg_id:577486 Sphingomonas yabuuchiae str. A1-18; GTC 868 NR_02
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FJ382671.1 gg_id:590918 Potentially pathogenic and air stem cell transplant summer hospital shower water BM FJ3826
X89903.1 gg_id:31767 Sphingomonas sp. str. PC5.2 (bright yel group 1 colony type) PC5.2 X89903
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GQ389119.1 gg_id:584563 quality deterioration drinking system during serious red water outbreak clone S72 GQ389
FJ525442.1 gg_id:536944 Chlorophyll-containing deep-sea deep sea water 500 m depth clone Pac500QII-10 FJ5254
FJ382620.1 gg_id:589104 Potentially pathogenic and air stem cell transplant summer hospital shower water BM FJ3826
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Y09638.1 gg_id:6244 Sphingomonas mali str. IFO 10550-T Y09638
AF128870.1 gg_id:105477 Sphingomonas sp. str. B26 AF128
EU834796.1 gg_id:579177 Microbiology Aerobic Phosphorus Removal Process aerobic phosphorus-removing se EU834
GQ161989.1 gg_id:588949 Sphingomonas sp. str. S8-3 GQ161
EU133552.1 gg_id:236646 tallgrass prarie soil clone FFCH3058 EU1335
EU730903.1 gg_id:356103 Sphingomonas melonis str. 150 EU730
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AY453855.1 gg_id:105178 Sphingomonas phyllosphaerae str. FA2 AY4538
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EU071479.1 gg_id:251041 and resistance microorganisms European clean room ESTEC HYDRA facility clone EHF EU0714
DQ532204.1 gg_id:162721 Geographical Variation Microbial Burden and Spacecraft Assembly Johnson Space Ce DQ532
AM697187.1 gg_id:226138 sequencing libraries indoor dust clone BF0001C028 AM697

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EU509065.1 gg_id:266499 cecal contents Mus musculus strain C57BL/6J; MD7 clone MD7_aap43a04 EU5090
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FJ382733.1 gg_id:509865 Potentially pathogenic and air stem cell transplant summer hospital shower water BM FJ3827
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EF028689.1 gg_id:197687 Sphingomonas sp. str. BCL-98 EF0286
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EU464953.1 gg_id:292322 African elephant feces clone AFYEL_aaj65g04 EU4649
AJ871278.1 gg_id:114482 Sphingomonas sp. str. DS3-1 AJ8712
EU816422.1 gg_id:346880 Sphingopyxis sp. str. Geo48 EU8164
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EF523589.1 gg_id:222242 Sphingopyxis sp. str. Ds26 EF5235
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EU075217.1 gg_id:242041 Sphingopyxis soli DCY34 EU0752
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EU143356.1 gg_id:248554 Sphingopyxis sp. str. J24 EU1433
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AB245343.1 gg_id:147540 Sphingopyxis ginsengisoli str. Gsoil 250 AB2453
AB219940.1 gg_id:132488 Sphingosinicella microcystinivorans str. MDB2 AB2199
EU703332.1 gg_id:384499 lakes on Eastern Tin Plateau freshwater lake altitude 4298 masl clone XZTSH2 EU7033
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DQ087403.1 gg_id:128144 Alisphingobium solum str. KSL-125 DQ087
AJ620201.1 gg_id:99158 Sphingomonas sp. str. PlS5 AJ6202
EU403958.1 gg_id:305626 Microbial septic tank effluent and soil biozone infiltrative surface clean water clone (EU4039
AJ303009.1 gg_id:52593 Novosphingobium lentum str. MT1 AJ3030
EU016180.1 gg_id:239608 Sphingomonas sp. str. S11-S-9 EU0161
DQ366009.1 gg_id:152371 Dominant soils Marble Point and Wright Valley Victoria Land Antarctica soil clone 20 DQ366
EU937972.1 gg_id:311613 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU9379
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GQ389106.1 gg_id:565345 quality deterioration drinking system during serious red water outbreak clone S60 GQ389
FJ675209.1 gg_id:525875 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6752
DQ823172.1 gg_id:171409 Techniques Microbial Studies Oregon Caves National Monument Quad 3 on trail clo DQ823
FJ478653.1 gg_id:582012 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4786
GQ389029.1 gg_id:570614 quality deterioration drinking system during serious red water outbreak clone J76 GQ389
AF170744.1 gg_id:6263 Sphingomonas sp. str. QSSC5-6 AF1707
FN397673.1 gg_id:426622 Blastomonas sp. str. AMV5 FN3976
D16149.1 gg_id:6203 Sphingomonas sp str. SYK6 D16149
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EU573097.1 gg_id:278193 and Dependent Microbial Oil Reservoir North Sea Ekofisk oil field Norwegian sector c EU573
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EU642848.1 gg_id:278075 Sandarakinorhabdus sp. JC2488 str. JC2488; IMSNU 14142 EU6428
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AF498741.1 gg_id:58089 soil isolate Ellin359 AF4987
GQ264264.1 gg_id:555926 Structure Simulated Level Waste Site simulated level waste site clone WW1_b69 GQ264
AY913375.1 gg_id:113174 forest soil clone DUNssu168 (-7A) (OTU#169) AY9133
AB081581.3 gg_id:74124 Rhizobiales str. A48 AB0815

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EF516517.1 gg_id:225609 grassland soil clone FCPP475 EF5165
DQ395957.1 gg_id:156154 deep-sea octacoral clone ctg_CGOGA24 DQ395
DQ659210.1 gg_id:223869 acid mine drainage clone DBS-Clone35 DQ659
AY921822.1 gg_id:114227 farm soil clone AKYG1082 AY9218
GQ346863.1 gg_id:544302 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ346
GQ340112.1 gg_id:582308 Prokaryotic Marathonas Reservoir Greece water column clone B07-10-BAC GQ340
EU801969.1 gg_id:315241 It's all ranking aquatic Chesapeake Bay MD clone 3C003352 EU801
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EF018958.1 gg_id:210397 aspen rhizosphere clone Amb_16S_1472 EF0189
EU907883.1 gg_id:346206 Identification hexahydro-135-trinitro-135-triazine (RDX)-degrading microorganisms v EU907
EF019039.1 gg_id:209880 aspen rhizosphere clone Amb_16S_1594 EF0190
AY234503.1 gg_id:111561 soil isolate Ellin5086 AY2345
FJ466360.1 gg_id:542729 and carbon monoxide-oxidizing across successional on recent Hawaiian 1959 Kilauea FJ4663
DQ001641.1 gg_id:158741 Microbial seeps asphalt seep clone E101-25 DQ001
EF032743.1 gg_id:177583 cyanobacterial mat in lava cave clone HAVOmat80 EF0327
AB198460.1 gg_id:135284 termite gut wall fraction clone RsW01-024 AB1984
FJ371976.1 gg_id:366881 feces adult twins and mothers; TS8 clone TS8_a04d02 FJ3719
FJ683333.1 gg_id:518552 Synecology primary and secondary habitats Escherichia coli O157:H7 fecal MARC bee FJ6833
AY916278.1 gg_id:113865 human rectum biopsy clone V326 AY9162
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EU460122.1 gg_id:296361 mammals their gut microbes eastern black and white colobus feces clone COL_aai15 EU460
EU767216.1 gg_id:348521 Pervasive effects antibiotic on human gut microbiota deep sequencing fecal clone C2 EU767
EU939387.1 gg_id:325850 Fecal wild pygmy loris feces clone WF16S_04 EU9393
GQ898126.1 gg_id:581652 Diets enriched fructans and/or soy protein can alter dominant fecal humans but do GQ898
GQ448745.1 gg_id:553416 Microbiome neonatal calves: shedding microorganisms fecal clone calf784_10wks_g GQ448
EU472130.1 gg_id:288432 francois langur feces clone FL_1aaa01a11 EU472
FJ683799.1 gg_id:518042 Synecology primary and secondary habitats Escherichia coli O157:H7 fecal MARC bee FJ6837
GQ396986.1 gg_id:538635 Functional unvegetated perhumid recently-deglaciated soils do not correlate shifts GQ396
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EF516476.1 gg_id:225263 grassland soil clone FCPP524 EF5164
FJ004770.1 gg_id:548087 Acidobacteria former agricultural bulk soil clone 32C1B FJ0047
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CU925843.1 gg_id:538971 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU9258
FJ517070.1 gg_id:540307 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled water c FJ5170
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EU871355.1 gg_id:343481 Genetic Bovine (Bos taurus) Rumen Thailand clone CF45 EU8713
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EU463527.1 gg_id:297362 domesticated horse feces clone horsem_aai95e02 EU463
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FJ202882.1 gg_id:345859 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2028
EU369125.1 gg_id:324513 libraries oyster shell clone MBIOS-09 EU369
FJ792144.1 gg_id:536627 Unexpectedly archaeal species shift between rare and dominant over thousand year FJ7921
EF092222.1 gg_id:265205 Persistent Axinella corrugata Rhizobiales clone B255_A3 EF0922
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EU328098.1 gg_id:278145 dynamics during bioremediation crude oil contaminated moderate saline soil clone Y EU328
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FJ497527.1 gg_id:537810 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4975
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AY945893.1 gg_id:149766 Thauera and Azoarcus functionally important genera denitrifying quinoline-removal AY9458
AM930304.1 gg_id:535965 gradual cooling stage mass and its spatial composting clone SMG103 AM930
GQ028948.1 gg_id:381203 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw85 GQ028
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EF648097.1 gg_id:237878 dynamics during produced water treatment aerobic activated sludge clone HB108 EF648
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CU466788.1 gg_id:342450 Evry wastewater treatment plant anoxic basin clone CU466
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EF516974.1 gg_id:223237 grassland soil clone FCPP392 EF5169
EF032738.1 gg_id:198250 cyanobacterial mat in lava cave clone HAVOmat41 EF0327
AJ536878.1 gg_id:78776 uranium mining waste pile soil sample clone JG30-KF-AS50 AJ5368
EU982844.1 gg_id:335424 Rhodobium sp. str. 6 EU982
EU104244.1 gg_id:325439 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104

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AY957890.1 gg_id:114903 drinking water biofilm clone B1NR70D5
DQ889887.1 gg_id:166172 natural product producing octocoral Erythropodium caribaeorum clone EC154
DQ856551.1 gg_id:255188 intestinal microflora Chinese mitten crab (Eriocheir sinensis) clone C3M
NC_010645.1 gg_id:281797 Bordetella avium str. 197N
AY946288.1 gg_id:114767 Achromobacter xylosoxidans 48
EU862292.1 gg_id:353739 Achromobacter ruhlandii str. jql-9
EF465456.1 gg_id:215887 Achromobacter xylosoxidans SF2
AY081976.1 gg_id:68623 atrazine-catabolizing microbial presence methanol clone KRA30+05
DQ414679.1 gg_id:207934 Achromobacter xylosoxidans
EU442188.1 gg_id:290452 Achromobacter xylosoxidans str. IdW
EU082170.1 gg_id:249176 Bordetella hinzii str. AU12584
AY081977.1 gg_id:68624 atrazine-catabolizing microbial presence methanol clone KRA30+06A
EF102817.1 gg_id:247706 Achromobacter xylosoxidans str. N-5-20
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EU308119.1 gg_id:240269 Achromobacter xylosoxidans str. WT1
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AY468369.1 gg_id:100569 Achromobacter xylosoxidans str. DTgA
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FJ810080.1 gg_id:536139 Achromobacter denitrificans str. 22426
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EU434544.1 gg_id:268968 Achromobacter xylosoxidans str. b62
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AF205370.1 gg_id:49793 Achromobacter ruhlandii str. QSSC3-1
EU877076.1 gg_id:311931 Achromobacter xylosoxidans str. EC3_3502

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AY753652.1 gg_id:107787 Achromobacter xylosoxidans str. VH070857 AY7536
AJ277798.1 gg_id:38214 Bordetella trematum str. DSM 11334 (T) AJ2777
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EF035138.1 gg_id:182422 Achromobacter xylosoxidans str. L2-2 EF0351
FN433031.1 gg_id:537770 Alcaligenes faecalis CCM5B FN4330
EU827475.1 gg_id:345476 Achromobacter xylosoxidans str. TFD47 EU8274
EU195864.1 gg_id:250159 Alcaligenes sp. str. ACS1 EU1958
EU221508.1 gg_id:332634 Trachelomonas armata str. ACOI 1323 EU2215
AY221606.1 gg_id:89748 metal and hydrocarbon contaminated soil clone HN4-32 AY2216
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EU652103.1 gg_id:273424 Achromobacter xylosoxidans str. ybc-11 EU6521
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FJ224081.1 gg_id:344427 Achromobacter denitrificans str. BMB-N6 FJ2240
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EU082160.1 gg_id:242404 Bordetella avium str. AU10563 EU0821
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FJ658470.1 gg_id:512527 Metalworking Fluids and Aerosols Metallurgy Plant metalworking aerosol clone Winte FJ6584
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AB188125.1 gg_id:132621 Alcaligenes latus str. IAM 12599 AB188
EF032746.1 gg_id:185782 cyanobacterial mat in lava cave clone HAVOmat12 EF0327
AY435511.1 gg_id:107880 propane-sparged ground water well C4 clone MAFB-C4-17 AY4355
FJ535109.1 gg_id:557756 microbial colonizing walls Karstic Slovenia cave wall biofilm clone 3PJM33 FJ5351
AB487742.1 gg_id:589854 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB487
AB486168.1 gg_id:579191 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSBARC AB486
AJ867848.1 gg_id:169526 Leptothrix sp. B2 AJ8678
AY493948.1 gg_id:136596 Fingerprinting aggregates soil clone 11 AY4939
AB378584.1 gg_id:260565 dominate denitrification nitrate- succinate-ammended stable istope probing 13C fra AB3785
AY466115.1 gg_id:101912 Bordetella holmesii G6128 AY466
EU082165.1 gg_id:244327 Bordetella petrii str. AU11495 EU082
DQ132877.1 gg_id:137970 Bordetella sp. str. QJ9-6 DQ132
EU082157.1 gg_id:251205 Bordetella sp. str. AU9919 EU082
DQ256466.1 gg_id:142658 Bordetella sp. str. 1-3 DQ256
EU835434.1 gg_id:346019 structure and quorum sensing reverse osmosis RO membrane biofilm clone 2B04 EU835
EU643523.1 gg_id:280840 Bordetella bronchiseptica str. Bb5 EU6435
AB255107.1 gg_id:171796 corroded concrete clone SC-127 AB255
EU603444.1 gg_id:263781 Bordetella petrii str. PTG4-17 EU6034
AJ249861.1 gg_id:21781 Bordetella petrii str. DSM 12804 Se-1111R AJ2498
EU082151.1 gg_id:240317 Bordetella sp. str. AU7206 EU082
AB039335.1 gg_id:37958 Bordetella sp. str. M1-6 AB0393
E06073.1 gg_id:25458 Bordetella bronchiseptica E06073
AY594190.1 gg_id:126199 Bordetella ansorpii str. SMC-8986 AY5941
AF142326.1 gg_id:6889 Bordetella pertussis str. Tohama; CIP 81.32 AF1423
AF366577.1 gg_id:49946 Bordetella parapertussis str. B24 AF3665
GQ480840.1 gg_id:541437 Bordetella petrii str. B1 GQ480
EU000383.1 gg_id:250883 Bordetella sp. str. BK-52 EU000
FJ969842.3 gg_id:445459 Bordetella parapertussis str. HT200 FJ9698
AJ005447.1 gg_id:6845 Alcaligenes defragrans str. 54Pin AJ0054
AB195160.1 gg_id:108771 Alcaligenes defragrans str. PD-16 AB195
Y12639.1 gg_id:6856 Denitrobacter permanens Y12639
EF205259.1 gg_id:189889 Alcaligenes sp. 'TJOPY2 (-I)' str. TJOPY2 (-I) EF2052
AF227163.1 gg_id:71251 Oligella urethralis str. ATCC17960 AF2271
AJ251912.1 gg_id:35834 Oligella ureolytica str. LMG 6519 AJ2519
AF190911.1 gg_id:6840 Pelistega europaea str. LMG15725 AF1909
FN421876.1 gg_id:594220 proteogenomics physiology phyllosphere clover clone 2_F01 FN4218
EU583723.1 gg_id:263890 Pigmentiphaga sp. str. JSM 061001 EU5837
EU170477.1 gg_id:238547 Pigmentiphaga sp. str. Zn-d-2 EU1704
AB300989.1 gg_id:272661 Sutterella parvirubra str. YIT 11816 (= JCM 14724 = DSM 19 AB3009
GQ033291.1 gg_id:455791 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw: GQ033

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EF400872.1 gg_id:213646 human fecal clone SJTU_D_02_65 EF4008
AJ832129.1 gg_id:181656 Sutterella morbirenis str. CCUG 49335 AJ8321
EU532008.1 gg_id:271525 Predominant role host genetics controlling gut microbiota FMF remission stool clone EU532
EU009806.1 gg_id:254552 intestinal microbiota domestic and wild turkeys turkey cecum clone WTB_D10 EU0098
EU009787.1 gg_id:245510 intestinal microbiota domestic and wild turkeys turkey cecum clone DTB_T27 EU0097
AB491209.1 gg_id:547221 Parasutterella sp. str. YIT 12071 AB491
EU767287.1 gg_id:339461 Pervasive effects antibiotic on human gut microbiota deep sequencing fecal clone C2 EU767
GQ011110.1 gg_id:498556 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ011
AJ566849.1 gg_id:104793 Sutterella stercoricanis str. CCUG 47620 AJ5668
GQ031994.1 gg_id:445073 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ031
EU767511.1 gg_id:322193 Pervasive effects antibiotic on human gut microbiota deep sequencing fecal clone C2 EU767
EU763570.1 gg_id:319660 Pervasive effects antibiotic on human gut microbiota deep sequencing fecal clone A4 EU763
EF071208.1 gg_id:176809 library survey and quantitative measurement major mucosa- IBD human colonic muc EF0712
AY981032.1 gg_id:126136 human rectum mucosal biopsy clone V241 AY9810
FJ368987.1 gg_id:358132 feces adult twins and mothers; TS4 clone TS4_a02h09 FJ3689
EU767098.1 gg_id:345647 Pervasive effects antibiotic on human gut microbiota deep sequencing fecal clone C1 EU767
DQ795867.1 gg_id:196501 human fecal clone RL245_aai83f07 DQ795
NR_024915.1 gg_id:540774 Taylorella asinigenitalis str. UCD-1; ATCC 700933 NR_024
AB200410.1 gg_id:112165 Taylorella equigenitalis str. Aus4 AB2004
AJ864470.1 gg_id:109028 Tetrathiobacter kashmirae str. WT001 AJ8644
AJ864472.1 gg_id:109030 Tetrathiobacter kashmirae str. 445C AJ8644
AJ864474.1 gg_id:109032 Tetrathiobacter kashmirae str. WPT AJ8644
AJ864473.1 gg_id:109031 Tetrathiobacter kashmirae str. WGT AJ8644
AJ864471.1 gg_id:109029 Tetrathiobacter kashmirae str. 445A AJ8644
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AY880023.1 gg_id:111634 Alcaligenes monasteriensis str. DPN7 AY8800
X86584.1 gg_id:6881 Alcaligenes sp str. 05-51 X86584
AM944735.1 gg_id:323924 Advenella incenata 4 AM944
DQ469586.1 gg_id:157497 Alcaligenes monasteriensis str. Ac2-VH DQ469
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GQ203113.1 gg_id:558099 Ochrobactrum pseudogrignonense str. 21-6PIN GQ203
AM944734.1 gg_id:348042 Advenella incenata str. CCUG 45225 AM944
EU447156.1 gg_id:273668 Tetrathiobacter kashmirensis str. CA1 EU447
AY622228.1 gg_id:106744 subsurface soil clone C-CG17 AY6222
FJ477679.1 gg_id:511551 Bordetella sp. HU-1 str. 1 FJ4776
U82826.1 gg_id:6846 Alcaligenes sp. str. NKNTAU U82826
AY792261.1 gg_id:120103 Humic Lake clone CrystalBog1G5 AY7922
EU801712.1 gg_id:309818 It's all ranking aquatic Chesapeake Bay MD clone 3C003072 EU801
EU703295.1 gg_id:466157 lakes on Eastern Tin Plateau polysaline lake altitude 3204 masl clone XZQH78 EU703

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DQ346505.1 gg_id:152918 during composting libraries compost clone 2B21 DQ346
EF603941.1 gg_id:230030 Salmonella typhimurium Exploits Inflammation Compete Intestinal Microbiota mouse EF6039
FM253637.1 gg_id:315961 Rock gold mine unusually microbial biofilm clone B01-2 FM253
EF097495.1 gg_id:194862 mouse cecum clone SWPT19_aaa03d11 EF0974
AY921968.1 gg_id:112764 farm soil clone AKYG1661 AY9219
AY986009.1 gg_id:127098 human stool clone D320 AY9860
EF405056.1 gg_id:215097 human fecal clone SJTU_G_07_39 EF4050
GQ254282.1 gg_id:540496 Bordetella sp. str. H12 GQ254
AY662006.1 gg_id:137438 uranium-contaminated groundwater clone 015B-B03 AY6620
DQ824015.1 gg_id:189076 human fecal clone RL204_aaj58b08 DQ824
AB246799.2 gg_id:150109 Alcaligenes sp. str. R-3 AB2467
EU510311.1 gg_id:273356 cecal contents Mus musculus strain C57BL/6J; WD3 clone WD3_aak04e08 EU5103
AY792260.1 gg_id:119586 Humic Lake clone CrystalBog1E9 AY7922
DQ905687.1 gg_id:165322 human fecal clone 014C-G4 DQ905
FJ973586.1 gg_id:384641 Periodic sulfide irruptions impact microbial structure and hypersaline lake water colu FJ9735
FJ477590.1 gg_id:586368 gut symbionts are tightly linked herbivory ants whole extracted single ant clone 86a FJ4775
AJ005449.1 gg_id:6852 Alcaligenes defragrans str. 62Car AJ0054
EF632938.1 gg_id:253486 Unique microbial contrasting aquatic altitude Andean Altiplano (northern Chile) fresh EF6329
AB259961.1 gg_id:160887 Burkholderia sp. str. Yoon-02 AB2599
AJ575701.1 gg_id:95883 lake water enrichment clone NE78 AJ5757
FJ672653.1 gg_id:528864 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6726
DQ466075.1 gg_id:160810 Pusillimonas terrae DQ466
EF405240.1 gg_id:208210 human fecal clone SJTU_G_04_82 EF4052
AJ786800.1 gg_id:108077 Alcaligenes sp. R-21939 AJ7868
EU888003.1 gg_id:303810 Microbial structure hydrolytic stage two-stage anaerobic digestion grass silage leach EU8880
AY238499.1 gg_id:92793 Alcaligenes sp. str. 92 AY2384
FJ901002.1 gg_id:547998 Polymorphism microbial reservoirs water cut stage oil field clone G-51 FJ9010
AY993322.1 gg_id:130981 mouse cecum clone C21_o12 AY9933
EU887771.1 gg_id:339734 and RFLP anaerobes main Nisargruna biogas plant anaerobic digester clone 2SN EU8877
FJ675537.1 gg_id:526614 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6755
EF096018.1 gg_id:189188 mouse cecum clone lean2_aaa02b05 EF0960
FJ960346.1 gg_id:541465 bottlenose dolphin upper respiratory tract blowhole clone D9A4_244 FJ9603
EF175378.1 gg_id:174707 Castellaniella sp. 7.5A2 EF1753
FJ880692.1 gg_id:404082 Diets enriched oat or temporally and alter fecal rat feces animal fed wheat bran diet; FJ8806
AJ318125.1 gg_id:26755 Waste-gas biofilter clone BIfciii38 AJ3181
EU511968.1 gg_id:273822 cecal contents Mus musculus strain C57BL/6J; WD2 clone WD2_aak12a12 EU5119
L29265.1 gg_id:6860 Blastocrithidia culicis Endosymbiont L29265
EF672088.1 gg_id:256466 Pusillimonas sp. str. DCY25T EF6720
DQ799851.1 gg_id:187938 human fecal clone RL182_aah34h10 DQ799
EF471693.1 gg_id:258429 Fast Screening and Estuarine Operon Libraries whole surface water Chesapeake Bay EF4716

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AY672759.1 gg_id:132213 Alcaligenes sp. 80 AY6727
AM982630.1 gg_id:279278 microbial and slurry pig saw dust spent bedding clone PISD-AMH11 AM982
EF442019.1 gg_id:214423 Bordetella petrii str. PLLA-3 EF4420
AY983565.1 gg_id:119964 human descending colon mucosal biopsy clone NN45 AY9835
FJ755942.1 gg_id:512238 Alcaligenes sp. str. RCT1 FJ7559
AY770946.1 gg_id:108314 water injection well Dagang oilfield clone W6 AY7709
FJ477560.1 gg_id:554733 gut symbionts are tightly linked herbivory ants whole extracted mid- and hind-gut sin FJ4775
FJ959932.1 gg_id:580692 bottlenose dolphin upper respiratory tract blowhole clone D8A8_326 FJ9599
DQ799318.1 gg_id:187226 human fecal clone RL205_aaj20g09 DQ799
DQ469227.1 gg_id:158061 Microbial and ecology water Zhong Tiaoshan mine Shanxi province China AMD clone DQ469
AY963392.1 gg_id:154501 structures and correlated ecological factors soils two evergreen broad-leaved forests AY9633
AY131213.1 gg_id:59999 Kerstersia gyiorum str. LMG 5906 AY1312
EU528222.1 gg_id:282571 Seasonal Dynamics Mudflat Mouth Major Kentucky Lake Reservoir Tributary sedime EU5282
AB487629.1 gg_id:562108 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB4876
EF097368.1 gg_id:177104 mouse cecum clone SWPT19_aaa01f07 EF0973
GQ422442.1 gg_id:586902 Pusillimonas terrae str. SAG3 GQ422
DQ451484.1 gg_id:156890 forest soil clone FAC45 DQ451
FJ768950.1 gg_id:537405 Alcaligenes sp. str. WY200811 FJ7689
EF999384.1 gg_id:250119 Vertical and Methanotrophs Estuarine Sediments Pearl River Estuary sediments 22cn EF9993
AJ867907.1 gg_id:144374 Fluctuations Significantly Influence Microbial Present Nival Lakes lake water clone B4 AJ8679
EF471697.1 gg_id:277235 Fast Screening and Estuarine Operon Libraries whole surface water Chesapeake Bay EF4716
GQ254283.1 gg_id:574340 Castellaniella sp. str. H13 GQ254
FJ477650.1 gg_id:584565 gut symbionts are tightly linked herbivory ants whole extracted single ant clone 96j FJ4776
FJ152724.1 gg_id:317726 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX5A_ FJ1527
EF520469.1 gg_id:228829 structure lakes: what controls ? acid-impacted lake clone ADK-HDe02-53 EF5204
FM873876.1 gg_id:368602 occupant source house mattress dust clone MB02C10 FM873
EF471233.1 gg_id:220259 Alcaligenes sp. str. BBTR16 EF4712
X92415.1 gg_id:6857 Alcaligenes sp. str. L6 X92415
DQ802483.1 gg_id:196880 human fecal clone RL188_aah18d02 DQ802
DQ799440.1 gg_id:184082 human fecal clone RL305aal86c09 DQ799
AJ277742.1 gg_id:22985 Brackiella oedipodis str. LMG 1945 R8846 AJ2777
AF232922.1 gg_id:22530 United Kingdom: Montserrat geothermal springs clone MS8 MS8 AF2329
EU133883.1 gg_id:237924 tallgrass prarie soil clone FFCH2283 EU1338
EF602797.1 gg_id:228235 Salmonella typhimurium Exploits Inflammation Compete Intestinal Microbiota mouse EF6027
GU138383.1 gg_id:585406 Achromobacter sp. str. THG 42 GU138
DQ346507.1 gg_id:153290 during composting libraries compost clone 2B23 DQ346
AY921769.1 gg_id:114410 farm soil clone AKYG1037 AY9217
AY792262.1 gg_id:123064 Humic Lake clone CrystalBog5E7 AY7922
EU800906.1 gg_id:330902 It's all ranking aquatic Delaware Bay NJ clone 2C229148 EU8009

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GQ102941.1 gg_id:415230 Topographical and Temporal Human Skin Microbiome skin inguinal crease clone nbv GQ102
GQ114946.1 gg_id:373715 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw685d0 GQ114
GQ107279.1 gg_id:429315 Topographical and Temporal Human Skin Microbiome skin popliteal fossa clone nbw GQ107
DQ256345.1 gg_id:143304 subsurface water clone EV818BHEB5102702SAS26 DQ256
GQ094491.1 gg_id:468438 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ094
GQ102742.1 gg_id:438634 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw(GQ102

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GQ094349.1 gg_id:453897 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ094

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GQ105736.1 gg_id:373767 Topographical and Temporal Human Skin Microbiome skin axillary vault clone nbw5 GQ105
EF653582.1 gg_id:240768 water s Dong Tinghu clone DTH12 EF6535

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GQ085424.1 gg_id:422806 Topographical and Temporal Human Skin Microbiome skin elbow clone nbw275e05(GQ085
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GQ090536.1 gg_id:427225 Topographical and Temporal Human Skin Microbiome skin popliteal fossa clone nbw GQ090
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GQ085331.1 gg_id:444595 Topographical and Temporal Human Skin Microbiome skin elbow clone nbw274b02(GQ085

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GQ105639.1 gg_id:415200 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw! GQ105
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GQ094529.1 gg_id:413739 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ094
GQ097754.1 gg_id:433206 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw456c02 GQ097
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GQ104249.1 gg_id:443362 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ104
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GQ069930.1 gg_id:396520 Topographical and Temporal Human Skin Microbiome skin buttock clone nbw243e0 GQ069

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AB365791.1 gg_id:257466 Burkholderia sp. str. WR43 AB365
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EF178444.1 gg_id:188143 Burkholderia cepacia str. 2Pe22 EF1784
EU477812.1 gg_id:258305 and validation prototype -based taxonomic microarray members genus Burkholderia EU477
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CP001026.1 gg_id:263782 Burkholderia ambifaria str. MC40-6 CP001
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AY741362.1 gg_id:106783 Burkholderia cepacia str. ATCC 53130 AY7413
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CP001504.2 gg_id:518847 Burkholderia glumae str. BGR1 CP0015
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AY268142.1 gg_id:91013 Burkholderia cepacia str. 1998019618 AY2681
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AM284970.1 gg_id:159660 Burkholderia nodosa R-22632 AM284
AB232335.1 gg_id:153688 Burkholderia sp. str. SFA3 AB2323
AJ551104.1 gg_id:94022 Burkholderia sp. wp26 AJ5511
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AB114607.1 gg_id:107108 Burkholderia cepacia str. CNR22 AB1146
AY792254.1 gg_id:119349 Humic Lake clone CrystalBog572G9 AY7922
AB288301.1 gg_id:211076 Burkholderia sp. str. BCB-57 AB2883
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EU477808.1 gg_id:261284 and validation prototype -based taxonomic microarray members genus Burkholderia EU4778
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EU399922.1 gg_id:266408 Burkholderia sp. str. CAF324 EU3999
AB537485.1 gg_id:586257 Burkholderia acidipaludis str. NBRC 101816 AB5374
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AJ251636.1 gg_id:28964 Scutellospora castanea strain BEG1 symbiont gigasporarum AJ2516
AJ251635.1 gg_id:50390 Scutellospora persica strain E09 symbiont AJ2516
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AF476077.1 gg_id:63963 Australicoccus grevilleae symbiont AF476
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AF476090.1 gg_id:63954 Nipaecoccus exocarpi symbiont AF476
AF476081.1 gg_id:63962 Cyphonococcus alpinus symbiont AF476
AF476079.1 gg_id:63964 Antonina pretiosa symbiont AF476
AF476092.1 gg_id:63949 Planococcus ficus symbiont AF476
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GQ284446.1 gg_id:541884 Chitinimonas taiwanensis str. THWCSN28 GQ284
DQ256728.1 gg_id:144420 Chitinimonas koreensis str. R2A43-10 DQ256
AY323827.2 gg_id:107650 Chitinimonas taiwanensis str. cf AY3238
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EU915714.1 gg_id:336731 Cupriavidus taiwanensis str. G EU9157
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EU024163.1 gg_id:251612 Cupriavidus gilardii str. CIP 105966 EU0241
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EU024156.1 gg_id:241006 Cupriavidus oxalaticus str. CIP 107389 EU0241
AB232327.1 gg_id:150729 Ralstonia sp. str. IFA3 AB2323
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EU024165.1 gg_id:239356 Cupriavidus pauculus str. CIP 105943 EU0241
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EU827495.1 gg_id:310258 Ralstonia eutropha str. JMP134 EU8274
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EF114432.1 gg_id:547037 Cupriavidus taiwanensis MSMB31 EF1144
GQ342296.1 gg_id:538911 Cupriavidus necator str. CG-T3 GQ342
AF067833.1 gg_id:7397 Ralstonia sp. str. TFD41 AF0678
AY583652.1 gg_id:102442 Ralstonia sp. str. 1-1 AY5836
AF500586.1 gg_id:66050 Wautersia respiraculi str. AU1618 AF5005
AY860251.1 gg_id:111512 Cupriavidus taiwanensis str. AUX0106 AY8602
AY571787.1 gg_id:100095 Ralstonia sp. str. 1-1 AY5717
AY860242.1 gg_id:111723 Cupriavidus respiraculi str. AU4630 AY8602
EU447184.1 gg_id:261584 Cupriavidus basilensis str. SE15 EU4471
D87999.1 gg_id:7386 Cupriavidus necator str. DSM 2839 D87999
AB007995.1 gg_id:58406 Wautersia eutropha str. P11 AB0079
EF114429.1 gg_id:545936 Cupriavidus gilardii MSMB32 EF1144
AF251026.1 gg_id:41961 Ralstonia sp. str. LD35 AF2510
AY860237.1 gg_id:111546 Cupriavidus respiraculi str. AU3775 AY8602
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AF312022.1 gg_id:73500 Wautersia basilensis str. DSM 11853 AF3120
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AY860227.1 gg_id:111839 Cupriavidus gilardii str. AU3713 AY8602
AY860234.1 gg_id:111890 Cupriavidus metallidurans str. AU4057 AY8602
AY860250.1 gg_id:111665 Ralstonia sp. str. AU5980 AY8602
AY303977.1 gg_id:88067 Wautersia taiwanensis str. MS1 AY3039
AB109778.1 gg_id:83846 Wautersia basilensis str. KF708 AB1097
NR_027607.1 gg_id:555836 Cupriavidus metallidurans CH34 str. CH34; ATCC 43123 NR_027
AF361022.1 gg_id:33253 Ralstonia sp. str. NAH8 AF3610
AB104447.1 gg_id:111989 Wautersia numadzuensis str. TE26 AB1044
AB271141.1 gg_id:168999 Cupriavidus gilardii str. MXC3-9 AB2711
EU827493.1 gg_id:325237 Cupriavidus respiraculi str. TFD48 EU8274
AF300325.2 gg_id:47468 Wautersia taiwanensis str. LMG 19425 AF3003
AF500587.1 gg_id:56268 Wautersia respiraculi str. AU3369 AF5005
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AB167180.1 gg_id:107898 Ralstonia sp. str. c310 AB1671
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AM905275.1 gg_id:327185 Cupriavidus taiwanensis JC4 AM905
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FJ823153.1 gg_id:563047 Lautropia sp. str. TeTO FJ8231

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GQ074337.1 gg_id:389328 Topographical and Temporal Human Skin Microbiome skin umbilicus clone nbw161g GQ074
AB366174.1 gg_id:239543 Limnobacter sp. str. KP1-19 AB366
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Y09879.1 gg_id:7183 Pandoraea norimbergensis Y09879
AY268174.1 gg_id:88183 Pandoraea norimbergensis str. CCUG 39188 AY268
EF596910.1 gg_id:230684 Pandoraea pnomenusa str. B-356 EF5969
AY268168.1 gg_id:88090 Pandoraea pnomenusa str. 2001008157 AY268
AB244466.1 gg_id:240824 Pandoraea pulmonicola str. C10-2 AB244
DQ837222.1 gg_id:255307 Archaeal and pristine National Park Spain Donana coastal aquifer clone 49S1_1B_3 DQ837
AY677092.1 gg_id:139269 Pandoraea sputorum TC84 AY677
AM921627.1 gg_id:238898 Pandoraea sputorum PhyCEm-128 AM921
AB118229.1 gg_id:105307 Pandoraea sp. PAMU-2.78 AB118
AY693821.1 gg_id:107349 anaerobic sludge clone EUB64 AY6938
AF139176.1 gg_id:72789 Pandoraea sputorum str. LMG 18819 AF139
AM110110.2 gg_id:139991 Polynucleobacter sp. str. MWH-Braz-FAM3C AM110
AF534432.1 gg_id:85148 Toolik Lake main station at 3 m depth clone TLM08/TLMdgge25a proteobacterium AF5344
AF534427.1 gg_id:86341 Toolik Lake main station at 3 m depth clone TLM03/TLMdgge33 proteobacterium AF5344
EF667903.1 gg_id:240040 Long-term maintenance species-specific microbiota basal metazoan Hydra epithelium EF6679
FM208180.1 gg_id:320042 Polynucleobacter sp. MWH-PoolGreenA3 FM208
AY947963.1 gg_id:142102 Synchrony and seasonality two temperate rivers USA: Massachusetts Ipswich River c AY9479
EU800885.1 gg_id:352838 It's all ranking aquatic Delaware Bay NJ clone 2C229123 EU8008
EU803597.1 gg_id:315886 It's all ranking aquatic Lake Gatun clone 5C231196 EU8035
AJ811013.1 gg_id:106508 Polynucleobacter sp. FC1 AJ8110
AJ270258.1 gg_id:48140 Ralstonia mannitolilytica str. LMG6866 AJ2702
EU024152.1 gg_id:251899 Ralstonia mannitolilytica str. CIP 107281 EU0241
DQ924950.1 gg_id:249852 Ralstonia solanacearum str. SL341 DQ924
AB024604.1 gg_id:50898 Ralstonia solanacearum MAFF 211266 AB0246
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DQ997838.1 gg_id:177991 Ralstonia pickettii str. HM-1 DQ997
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AB024608.1 gg_id:40253 Ralstonia solanacearum MAFF 301559 AB0246
FJ193860.1 gg_id:338003 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1938
FJ194034.1 gg_id:308094 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1940
EF562231.1 gg_id:249054 Subsurface microbial deep-granitic-fracture water colorado Henderson Molybdenum EF5622
FJ191398.1 gg_id:326241 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1913
EF207564.1 gg_id:210668 mangrove system sediment EF2075
FJ193764.1 gg_id:350437 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1937
AJ583178.1 gg_id:95147 radioactive site ground water clone S15D-MN15 AJ5831
FJ193746.1 gg_id:349604 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1937

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EF562200.1 gg_id:246230 Subsurface microbial deep-granitic-fracture water colorado Henderson Molybdenum EF5622
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FJ193974.1 gg_id:332274 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1939
FJ192332.1 gg_id:324427 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1923
NW_002196521.1 gg_id:341686 Ralstonia solanacearum str. MolK2 NW_00
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FJ193962.1 gg_id:323373 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1939
FJ193740.1 gg_id:308235 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1937
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DQ256353.1 gg_id:143763 subsurface water clone EV818BHEB5102702SAS31 DQ256
EU234261.1 gg_id:236043 river and un techniques downstream Wang Yang River receiving penicillin G producti EU2342
FJ625332.1 gg_id:511989 Influence lead contamination on boreal pine forest soil clone HF_M_29 FJ6253
CU466909.1 gg_id:313184 Evry wastewater treatment plant anoxic basin clone CU4669
AB089482.1 gg_id:96705 Derxia gummosa str. IAM 13946 AB0894
EU304284.1 gg_id:244827 Ralstonia sp. str. 22 EU3042
FJ516875.1 gg_id:558389 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled lower s FJ5168
CU927753.1 gg_id:558915 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU9277
FJ484886.1 gg_id:512042 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4848
EU133821.1 gg_id:253342 tallgrass prarie soil clone FFCH11971 EU1338
EU799756.1 gg_id:319986 It's all ranking aquatic Newport Harbour RI clone 1C227401 EU7997
FJ167462.1 gg_id:310487 microbiology granular sludge under autotrophic mixotrophic and heterotrophic denit FJ1674
DQ450176.1 gg_id:151384 (Uruguay) Atlantic coastal lagoon clone R7C70 DQ450
AJ853520.1 gg_id:110392 landfill leachate clone GZKB25 AJ8535
EU491796.1 gg_id:260119 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU4917
FJ612332.1 gg_id:545233 and Biogeochemistry Dongping Lake Ecosystems lake water clone DP10.1.39 FJ6123
EF654710.1 gg_id:236325 suspended sludge anammox reactor clone KIST-JJY041 EF6547
AB487015.1 gg_id:546434 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNIR0 AB4870
AY218566.1 gg_id:81664 penguin droppings sediments clone KD1-79 AY2185
EU703319.1 gg_id:412714 lakes on Eastern Tin Plateau polysaline lake altitude 3204 masl clone XZQH193 EU7033

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CU466912.1 gg_id:355680 Evry wastewater treatment plant anoxic basin clone CU466
AY921755.1 gg_id:113342 farm soil clone AKYH739 AY921
EU802154.1 gg_id:324381 It's all ranking aquatic Chesapeake Bay MD clone 3C003572 EU802
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AB240293.1 gg_id:141532 microbial structure Phragmites rhizosphere biofilm reed bed reactor laboratory clon AB240
FJ375442.1 gg_id:558588 T-RFLP powerful tool rapid assessment cells microbial fuel cell sheet carbon cloth ano FJ3754
FJ612382.1 gg_id:567560 and Biogeochemistry Dongping Lake Ecosystems lake water clone DP10.3.32 FJ6123
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FJ167492.1 gg_id:325718 microbiology granular sludge under autotrophic mixotrophic and heterotrophic denit FJ1674
AY570587.1 gg_id:108463 biodegraded Canadian oil reservoir clone PL-33B5 AY570
GQ010467.1 gg_id:499198 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ010
FJ230924.1 gg_id:326020 under Level Antibiotic Selection Pressures river water clone F2 FJ2309
AY509453.2 gg_id:103790 freshwater clone LiUU-5-131 AY509
FJ674610.1 gg_id:515098 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface FJ6746
EF516760.1 gg_id:225714 grassland soil clone FCPS725 EF5167
AB252925.1 gg_id:170130 Identification Shibayama lagoon Ishikawa Japan iron-oxidation biofilm clone 57 AB252
EU727457.1 gg_id:354510 Ralstonia sp. str. JOS55-1 EU727
GQ500805.1 gg_id:570888 Surveys Biofilms and Sediments Mammoth Cave Karst Aquifers Bio-Sep beads conta GQ500
GQ389057.1 gg_id:591556 quality deterioration drinking system during serious red water outbreak clone S13 GQ389
AM850114.1 gg_id:249856 Acidovorax avenae subsp. citrulli Med-H AM850
AM084006.1 gg_id:140023 Acidovorax sp. str. R-24608 AM084
EU434539.1 gg_id:264675 Acidovorax defluvii str. b322 EU434
EU434519.1 gg_id:275328 Acidovorax defluvii str. b224 EU434
EU434529.1 gg_id:272063 Acidovorax defluvii str. b336 EU434
EU434535.1 gg_id:276678 Acidovorax defluvii str. b256 EU434
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GQ284423.1 gg_id:552122 Acidovorax delafieldii str. THWCSN54 GQ284
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EU730927.1 gg_id:315092 Acidovorax facilis str. 228 EU730
EU304287.1 gg_id:251328 Acidovorax sp. str. 75 EU304
Y18616.1 gg_id:7043 Acidovorax defluvii str. BSB411 Y18616
AB487337.1 gg_id:553499 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNIR0 AB487
EU104261.1 gg_id:346700 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
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GU372409.1 gg_id:575562 Acidovorax sp. str. S7 GU372
GQ106011.1 gg_id:431652 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ106

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CU466871.1 gg_id:339277 Evry wastewater treatment plant anoxic basin clone CU466
FJ593725.1 gg_id:531432 midgut tropical caterpillars fed on diets larvae clone CLSM148 FJ5937
AY093698.1 gg_id:55345 Acidovorax sp. str. 'smarlab 133815' AY093
GQ091955.1 gg_id:414179 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ091
AB269774.1 gg_id:208383 Acidovorax delafieldii str. HYY0511-SK09 AB269
GQ107139.1 gg_id:397297 Topographical and Temporal Human Skin Microbiome skin popliteal fossa clone nbw GQ107
EU202950.1 gg_id:251020 Linking -situ microbial identity and functionality without bias groundwater naphthale EU202
GQ102018.1 gg_id:387956 Topographical and Temporal Human Skin Microbiome skin palm clone nbw502e06c1 GQ102
DQ264575.1 gg_id:143236 universal microarray broader than typical library when sampling subsurface groundv DQ264
AJ400840.1 gg_id:74685 Acidovorax wohlfahrtii str. B2/74 AJ4008
AB008429.1 gg_id:7063 Comamonas sp. str. MBIC3885 E6 AB008
AB167175.1 gg_id:107894 Acidovorax sp. str. c303 AB167
GQ107342.1 gg_id:440564 Topographical and Temporal Human Skin Microbiome skin popliteal fossa clone nbw GQ107
AF078762.1 gg_id:7047 Acidovorax avenae subsp. cattleyae str. NCPPB 961 subsp. AF078
GQ068971.1 gg_id:447274 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ068
DQ264434.1 gg_id:141042 universal microarray broader than typical library when sampling subsurface groundv DQ264
FJ560467.1 gg_id:546272 Acidovorax sp. str. SD2 FJ5604
AM084008.1 gg_id:140621 Acidovorax sp. str. R-24614 AM084
GQ022385.1 gg_id:487284 Topographical and Temporal Human Skin Microbiome skin axillary vault clone nbu31 GQ022
FM213427.1 gg_id:327583 Chemosynthetic Bivalvia species mud volcanoes Gulf Cadiz host gill tissue clone Mer FM213
GQ106875.1 gg_id:435090 Topographical and Temporal Human Skin Microbiome skin buttock clone nbw542a0 GQ106
GQ114179.1 gg_id:393759 Topographical and Temporal Human Skin Microbiome skin manubrium clone nbw59 GQ114
AF078760.1 gg_id:7052 Acidovorax konjaci str. ATCC 33996 AF078
AB076844.1 gg_id:28959 Acidovorax sp. str. OS-6 AB076
GQ098446.1 gg_id:400311 Topographical and Temporal Human Skin Microbiome skin umbilicus clone nbw559g GQ098
GQ099609.1 gg_id:451960 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw51 GQ099
AY837725.1 gg_id:141709 -Based Identification Midgut Field-Caught Anopheles gambiae Sensu Lato and . funes AY837
EU104224.1 gg_id:356016 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
GQ114282.1 gg_id:422535 Topographical and Temporal Human Skin Microbiome skin manubrium clone nbw59 GQ114
EU434475.1 gg_id:275733 Acidovorax defluvii str. b268 EU434
GQ099219.1 gg_id:447125 Topographical and Temporal Human Skin Microbiome skin interdigital web space clc GQ099
GQ102686.1 gg_id:413906 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ102
FJ497649.1 gg_id:585668 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4976
GQ104106.1 gg_id:447512 Topographical and Temporal Human Skin Microbiome skin glabella clone nbw563c0 GQ104
AF015487.1 gg_id:7077 Pseudomonas sp. str. P51 AF015
FJ658868.1 gg_id:568875 Metalworking Fluids and Aerosols Metallurgy Plant metalworking aerosol clone Sumn FJ6588
FJ477624.1 gg_id:552067 gut symbionts are tightly linked herbivory ants whole extracted single ant clone UAB- FJ4776
FJ477611.1 gg_id:578828 gut symbionts are tightly linked herbivory ants whole extracted mid- and hind-gut sin FJ4776
AF508114.1 gg_id:66230 Acidovorax avenae C1 AF508
FJ657748.1 gg_id:545408 Metalworking Fluids and Aerosols Metallurgy Plant metalworking aerosol clone Winte FJ6577

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FJ658952.1 gg_id:553262 Metalworking Fluids and Aerosols Metallurgy Plant metalworking aerosol clone Sumn FJ6589
FJ658390.1 gg_id:544123 Metalworking Fluids and Aerosols Metallurgy Plant metalworking aerosol clone Winte FJ6583
AB120964.1 gg_id:99037 Aquamonas fontana str. AQ10 AB120
DQ640723.1 gg_id:227889 full-scale EBPR sludge clone Skagen135 DQ640
AF526926.2 gg_id:62982 Mars Odyssey Orbiter and encapsulation facility clone T5-4 sp. AF5269
AY133071.1 gg_id:56377 TCE-contaminated site clone ccs221 AY1330
DQ521541.1 gg_id:193826 Antarctic lake ice cover clone ANTLV7_G03 DQ521
AY662013.1 gg_id:137452 uranium-contaminated groundwater clone 015C-G04 AY6620
AF526920.2 gg_id:62991 Mars Odyssey Orbiter and encapsulation facility clone T8-7 sp. AF5269
EU150266.1 gg_id:279514 acidobacteria and soil Niwot Ridge LTER clone Ni_F8P2 EU1502
NR_028713.1 gg_id:585432 Curvibacter delicatus str. 146; LMG 4328 NR_028
D14320.1 gg_id:6938 Brachymonas denitrificans str. -P1 D14320
EU434344.1 gg_id:271636 Brachymonas denitrificans str. a265 EU4343
EU434398.1 gg_id:265311 Brachymonas denitrificans str. a310 EU4343
AB286490.1 gg_id:208338 Reducing production and domination comamonadaceae reducing oxygen supply was AB2864
AJ318129.1 gg_id:21007 Waste-gas biofilter clone BIfdi44 AJ3181
AM420229.1 gg_id:239064 Noma subgingival plaque clone 601H06(oral) AM420
FJ982927.1 gg_id:460813 Comamonas sp. str. ST18 FJ9829
AB164432.1 gg_id:103333 Comamonas badia str. IAM 14839 AB1644
AJ576357.1 gg_id:107318 hindgut homogenate Pachnoda ephippiata larva clone PeH41 AJ5763
AF234720.1 gg_id:73090 sludge clone A6b AF2347
EU346912.1 gg_id:265983 Brachymonas sp. str. AImA4 EU3469
AB286456.1 gg_id:216049 Reducing production and domination comamonadaceae reducing oxygen supply was AB2864
FJ959768.1 gg_id:584835 bottlenose dolphin upper respiratory tract blowhole clone D8A8_102 FJ9597
AB286474.1 gg_id:214136 Reducing production and domination comamonadaceae reducing oxygen supply was AB2864
EU881354.1 gg_id:306436 (maize-sweet cropping field) Karst region maize-sweet potato cropland soil clone KM EU8813
AB286506.1 gg_id:211133 Reducing production and domination comamonadaceae reducing oxygen supply was AB2865
GQ010546.1 gg_id:499118 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ010
GQ110427.1 gg_id:447775 Topographical and Temporal Human Skin Microbiome skin back clone nbw659e10c1 GQ110
AM283038.1 gg_id:159681 Caldimonas sp. str. Han85 AM283
EU024140.1 gg_id:247295 Comamonas aquatica str. CIP 107986 EU0241
EU464974.1 gg_id:295063 African elephant feces clone AFYEL_aaj66a08 EU4649
DQ854969.1 gg_id:161904 Comamonas sp. str. GPTSA5 DQ854
EU464791.1 gg_id:289604 Hartmann's mountain zebra feces clone AFZEB_aaj70h03 EU4647
AF408327.1 gg_id:36914 Comamonas sp. clone KL-11-1-9 AF4083
EU841519.1 gg_id:337127 Comamonas aquatica str. 406 EU8415
DQ824524.1 gg_id:188323 human fecal clone RL185_aaj69c02 DQ824
AJ430344.1 gg_id:73664 Comamonas aquatica str. LMG 2370 AJ4303
AF233878.1 gg_id:40166 Comamonas denitrificans str. 2.99g AF2338
AF233879.1 gg_id:74723 Comamonas denitrificans str. 5.38g AF2338

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AJ430343.1 gg_id:66690 Comamonas terrigena str. LMG 1249	AJ4303
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EU024142.1 gg_id:240089 Comamonas terrigena str. LMG 5520	EU0241
AM184229.1 gg_id:160747 Comamonas terrigena str. WAB1888	AM184
NR_028719.1 gg_id:545968 Comamonas terrigena str. IMI 359870; ATCC 8461	NR_028
EU337122.1 gg_id:545663 Comamonas denitrificans str. 2B7	EU3371
GU454930.1 gg_id:524752 Understanding short-chain fatty acids accumulation enhanced alkaline digestion: kir	GU454
DQ824755.1 gg_id:189547 human fecal clone RL185_aaj72g08	DQ824
EU465136.1 gg_id:290700 African elephant feces clone AFYEL_aaj68e11	EU4651
FJ658617.1 gg_id:509630 Metalworking Fluids and Aerosols Metallurgy Plant metalworking aerosol clone Sumn	FJ6586
EU464722.1 gg_id:300641 Hartmann's mountain zebra feces clone AFZEB_aaj69f11	EU4647
AY566581.1 gg_id:108740 Comamonas sp. str. PG-08	AY5665
GQ179675.1 gg_id:465713 Heterogeneity vaginal microbial within individuals vagina clone VE3A04	GQ179
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EU465027.1 gg_id:296452 African elephant feces clone AFYEL_aaj66h05	EU4650
EU536017.1 gg_id:286592 profile human microbiota antecubital fossa (inner elbow) skin clone nbt161h12	EU5360
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EU465048.1 gg_id:302343 African elephant feces clone AFYEL_aaj67c02	EU4650
EU465099.1 gg_id:302440 African elephant feces clone AFYEL_aaj68a11	EU4650
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EU464843.1 gg_id:295086 Hartmann's mountain zebra feces clone AFZEB_aaj71g07	EU4648
EU465006.1 gg_id:293491 African elephant feces clone AFYEL_aaj66e11	EU4650
EU771834.1 gg_id:310863 African elephant feces clone AFYEL_aaj65h12	EU7718
AB076865.1 gg_id:42211 anaerobic sludge clone OS1L-7	AB0768
EU771819.1 gg_id:310322 African elephant feces clone AFYEL_aaj68f04	EU7718
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AY856846.1 gg_id:111198 Comamonas sp. str. 46	AY8568
EU464826.1 gg_id:287428 Hartmann's mountain zebra feces clone AFZEB_aaj71e07	EU4648
GQ389171.1 gg_id:571114 quality deterioration drinking system during serious red water outbreak clone D40	GQ389
EU771661.1 gg_id:334294 African elephant feces clone AFEL_aai29g11	EU7716
EU464941.1 gg_id:302302 African elephant feces clone AFYEL_aaj65e07	EU4649
EU465157.1 gg_id:292068 African elephant feces clone AFYEL_aaj68h04	EU4651
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DQ453128.1 gg_id:156701 Comamonas sp. str. Dant 3-8	DQ453

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AF501880.1 gg_id:71872 Comamonas testosteroni str. SMCC B0589 AF5018
EU467917.1 gg_id:289291 cheetah feces clone CE2_a08 EU4679
GU454975.1 gg_id:515113 Understanding short-chain fatty acids accumulation enhanced alkaline digestion: kir GU454
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GQ008024.1 gg_id:501642 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw: GQ008
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EU537205.1 gg_id:281659 profile human microbiota antecubital fossa (inner elbow) skin clone nbt68d07 EU5372
FJ893904.1 gg_id:540808 Matriptase-deficient mice exhibit ichthyotic selective shift skin microbiota B1E; ear pu FJ8939
AF532869.1 gg_id:91652 Comamonas sp. str. D1 AF5328
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EU465146.1 gg_id:297240 African elephant feces clone AFYEL_aaj68g02 EU4651
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DQ001153.1 gg_id:118400 Rhodobacter sphaeroides str. Il106 DQ001

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EU130955.1 gg_id:236660 Hydrogenophaga sp. BAC57 EU130
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DQ861288.1 gg_id:167427 Hylemonella gracilis ZL DQ861
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GU455142.1 gg_id:530933 Understanding short-chain fatty acids accumulation enhanced alkaline digestion: kir GU455

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AY792257.1 gg_id:124653 Humic Lake clone CrystalBog021H12 AY792
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EF667927.1 gg_id:245084 Long-term maintenance species-specific microbiota basal metazoan Hydra epitheliun EF6679
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AM501437.1 gg_id:240464 Pelomonas aquatica str. CCUG52631 AM501

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AM501440.1 gg_id:248405 Pelomonas puraquae str. CCUG52770 MG64 AM501
AF396932.1 gg_id:112368 Pseudomonas saccharophila str. PRO7 AF3969
DQ884343.1 gg_id:170667 Pseudomonas sp. str. JS-1 DQ884
AM039830.1 gg_id:133518 Polaromonas aquatica str. CCUG 39402 AM039
NC_008781.1 gg_id:194205 Polaromonas naphthalenivorans str. CJ2 NC_00
AAFQ02000005.1 gg_id:129807 Polaromonas sp. str. JS666 AAFQ0
DQ298150.1 gg_id:142773 -dependent and microbial assemblages silkworm midgut clone swm42 DQ298
AB464945.2 gg_id:323641 Altitudinal changes flora on Gulkana Glacier Alaska analyzed sequencing and -cycle c AB4649
AF361203.1 gg_id:53922 mesoeutrophic reservoir clone 16 AF3612
AY166684.1 gg_id:76891 Polaromonas naphthalenivorans str. CJ2 AY1666
U14585.1 gg_id:7005 Polaromonas vacuolata str. 34-P U14585
DQ341420.1 gg_id:150971 Acidovorax sp. str. Antarctic IS08 DQ341
AF407386.1 gg_id:33769 MCB-contaminated groundwater-treating reactor clone RB7C8 AF4073
AB252078.1 gg_id:152145 entrapped Puruogangri ice core clone PIC-D3 AB2520
EU801289.1 gg_id:323062 It's all ranking aquatic Chesapeake Bay MD clone 3C002543 EU8012
EU636027.1 gg_id:278933 Antarctic str. G020 EU6360
EU636024.1 gg_id:279595 Antarctic str. G076 EU6360
AY218637.1 gg_id:81748 penguin droppings sediments clone KD4-57 AY2186
EU636029.1 gg_id:278079 Antarctic str. G088 EU6360
FJ979854.1 gg_id:384326 Polaromonas sp. str. tsz24 FJ9798
FJ801199.1 gg_id:552577 Investigation Water -DGGE and Clone Library Zoige Alpine Wetland clone ZWB3-18 FJ8011
EU030285.1 gg_id:254221 Polaromonas jejuensis str. JS12-13 EU0302
EU801930.1 gg_id:355065 It's all ranking aquatic Chesapeake Bay MD clone 3C003311 EU8019
EF423322.1 gg_id:212990 Polaromonas sp. str. 1006 EF4233
EU703261.1 gg_id:448038 lakes on Eastern Tin Plateau oligosaline lake altitude 4718 masl clone XZNMC60 EU7032
DQ094183.1 gg_id:130292 Polaromonas sp. 'hydrogenovorans' str. hydrogenovorans DQ094
EU636028.1 gg_id:279079 Antarctic str. CC10 EU6360
AY188325.1 gg_id:79327 penguin droppings sediment clone KD7-99 AY1883
FM161402.1 gg_id:291214 Polaromonas sp. str. 01WB02.3-26 FM161
AB167243.1 gg_id:107951 str. c115 AB1672
AY168739.1 gg_id:75143 arsenite-oxidizing biofilm clone HC-3 AY1687
GU213397.1 gg_id:540184 Polaromonas sp. str. 190 GU213
EF516819.1 gg_id:224825 grassland soil clone FCPN433 EF5168
GU208230.1 gg_id:588612 Study on Sediment Dongping Lake sediment clone 3S4-1 GU208
AY921863.1 gg_id:113128 farm soil clone AKYH1084 AY9218
EU881341.1 gg_id:340905 (maize-sweet cropping field) Karst region maize-sweet potato cropland soil clone KM EU8813
DQ984611.1 gg_id:237973 LY oil-contaminated soil clone LYC180 DQ984
GQ079635.1 gg_id:390755 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw1232d(GQ079
AY439189.1 gg_id:104669 soil clone MeCl 24 AY4391
AJ532684.1 gg_id:76621 uranium mill tailings clone Gitt-KF-21 AJ5326

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DQ354711.1 gg_id:154792 subsurface water clone DR550SWSAEE7 DQ354
EF516224.1 gg_id:221803 grassland soil clone FCPU687 EF5162
AB487846.1 gg_id:591765 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB4878
FJ478510.1 gg_id:578098 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4785
DQ451482.1 gg_id:156693 forest soil clone FAC43 DQ451
EU979062.1 gg_id:317424 Effects intercropping Rhizobium inoculation on microbial biomass and rhizosphere b EU979
FM178814.1 gg_id:343929 Monitoring abundance and expression 'dehalococcoides' species chloroethene-redu FM178
NC_007908.1 gg_id:153264 Rhodoferax ferrireducens str. DSM 15236 NC_00
D16212.1 gg_id:6979 Rhodoferax fermentans str. FR3 D16212
X95837.1 gg_id:6986 situ visualization genetic natural microbial ecies strain T3 X95837
FJ517716.1 gg_id:513243 Disturbing epithelial homeostasis metazoan Hydra leads drastic changes microbiota e FJ5177
AY214178.1 gg_id:82365 benzene-contaminated groundwater clone ZZ9C20 AY2141
DQ628924.1 gg_id:162964 sequences Arctic glacier subglacial water ice or sediment clone SIB1 2B DQ628
DQ856540.1 gg_id:254818 intestinal microflora Chinese mitten crab (Eriocheir sinensis) clone C2M DQ856
AJ606333.1 gg_id:100951 Comamonadaceae SB1 AJ6063
DQ628925.1 gg_id:170859 sequences Arctic glacier subglacial water ice or sediment clone SIB2 1D DQ628
AB199580.1 gg_id:110990 river water Japan clone RVW-13 AB1995
EU907895.1 gg_id:315915 Identification hexahydro-135-trinitro-135-triazine (RDX)-degrading microorganisms v EU9078
AY218635.1 gg_id:81515 penguin droppings sediments clone KD4-45 AY2186
AF418942.1 gg_id:43083 USA: Colorado Fort collins Horsetooth Reservoir clone HTA10 AF4189
EF203797.1 gg_id:236168 Chlorochromatium aggregatum EF2037
AB154311.1 gg_id:104501 freshwater clone S9F-53 AB1543
EF451646.1 gg_id:216339 Rhodoferax sp. str. KAR16 EF4516
EF580989.1 gg_id:278249 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM7 EF5809
AJ421912.2 gg_id:27768 Germany:Elbe River clone Elb08 AJ4219
EF516373.1 gg_id:224371 grassland soil clone FCPT660 EF5163
AB237669.1 gg_id:140742 microorganisms deep subsurface groundwater sedimentary rock milieu clone HDBW AB2376
EU937983.1 gg_id:333026 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU9379
AJ867929.1 gg_id:142024 Fluctuations Significantly Influence Microbial Present Nival Lakes lake water clone D8 AJ8679
EU937978.1 gg_id:330894 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU9379
EU937973.1 gg_id:303742 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU9379
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GU233447.1 gg_id:516603 Rhodoferax antarcticus str. ANT.BR GU233
CU466889.1 gg_id:338390 Evry wastewater treatment plant anoxic basin clone CU466
AB475015.1 gg_id:566864 sequences located Engiadina Bassa Valley Switzerland iron containing freshwater so AB4750
AY214173.1 gg_id:81368 benzene-contaminated groundwater clone ZZ9C2 AY2141
AB478658.1 gg_id:534216 SSU sequences Alps Cadagno di Fuori submerged biofilm mats montane wetland clo AB4786
AB504936.1 gg_id:552998 structure microorganisms reddish-brown iron-rich snow clone oze05B33 AB5049
GQ340212.1 gg_id:552309 Prokaryotic Marathonas Reservoir Greece water column clone VE07-26-BAC GQ340
EF580979.1 gg_id:279495 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM1 EF5809

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AF435948.2 gg_id:75345 Rhodoferax ferrireducens str. T118 AF435
EF520458.1 gg_id:233033 structure lakes: what controls ? acid-impacted lake clone ADK-HDe02-15 EF5204
AJ422168.1 gg_id:34909 Germany:Spittelwasser River clone Spb249 AJ4221
AB475020.1 gg_id:534700 sequences located Engiadina Bassa Valley Switzerland iron containing freshwater so AB475
CU466882.1 gg_id:347539 Evry wastewater treatment plant anoxic basin clone CU466
EU340210.1 gg_id:248312 on Aquatic Macrophytes Geothermally Active Area Sedge Bay Yellowstone Lake WY EU340
AY013643.2 gg_id:23857 sample taken downstream landfill clone BVB72 landfill AY013
AY050600.1 gg_id:34176 MCB-contaminated groundwater-treating reactor clone GOUTB17 AY050
AF523029.1 gg_id:88367 natural mineral water clone G-5 AF523
EU104076.1 gg_id:340208 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
AM697182.1 gg_id:222656 sequencing libraries indoor dust clone BF0001C023 AM697
FM955857.1 gg_id:581777 Rhodoferax sp. str. Asd M2A1 FM955
FJ437765.1 gg_id:546951 Benthic okenone production Fayetteville New York Green Lake chemocline 20.5 m wa FJ4377
DQ628926.1 gg_id:171241 sequences Arctic glacier subglacial water ice or sediment clone SIB2 10G DQ628
GQ339123.1 gg_id:526119 Comprehensive Investigation on Iron Cycling Freshwater Seep ing Microscopy Cultiv GQ339
AJ867927.1 gg_id:144785 Fluctuations Significantly Influence Microbial Present Nival Lakes lake water clone D6 AJ8679
EF590026.2 gg_id:231559 changes structure sediments Songhua River contaminated nitrobenzene -DGGE cloni EF590
AJ421940.2 gg_id:20142 Germany:Elbe River clone Elb270 AJ4219
EU801480.1 gg_id:325766 It's all ranking aquatic Chesapeake Bay MD clone 3C002783 EU801
GQ389172.1 gg_id:587804 quality deterioration drinking system during serious red water outbreak clone D42 GQ389
AJ421927.2 gg_id:20305 Germany:Elbe River clone Elb154 AJ4219
AF523038.1 gg_id:90178 natural mineral water clone H-5 AF523
EU104187.1 gg_id:331535 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
DQ228371.1 gg_id:144642 microbial compositions two subglacial possible role microbes chemical weathering p DQ228
EU499697.1 gg_id:263522 Field-based laboratory stable isotope probing surveys identities both aerobic and an EU499
AB475018.1 gg_id:542491 sequences located Engiadina Bassa Valley Switzerland iron containing freshwater so AB475
EF520463.1 gg_id:231370 structure lakes: what controls ? acid-impacted lake clone ADK-MOh02-46 EF5204
EF590046.2 gg_id:230607 changes structure sediments Songhua River contaminated nitrobenzene -DGGE cloni EF590
EU636030.1 gg_id:278005 Antarctic str. G079 EU636
DQ227795.1 gg_id:142898 Rhodoferax sp. PIC-C33 DQ227
AJ421938.2 gg_id:18730 Germany:Elbe River clone Elb263 AJ4219
DQ664242.1 gg_id:172764 Rhodoferax sp. str. IMCC1723 DQ664
DQ228394.1 gg_id:143719 microbial compositions two subglacial possible role microbes chemical weathering p DQ228
DQ532326.1 gg_id:168650 Geographical Variation Microbial Burden and Spacecraft Assembly Kennedy Space C DQ532
AF407396.1 gg_id:50642 MCB-contaminated groundwater-treating reactor clone RB9C6 AF407
CU466916.1 gg_id:337770 Evry wastewater treatment plant anoxic basin clone CU466
AF407412.1 gg_id:44459 MCB-contaminated groundwater-treating reactor clone RB13C6 AF407
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DQ521494.1 gg_id:198759 Antarctic lake ice cover clone ANTLV1_G06 DQ521

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EU801168.1 gg_id:340336 It's all ranking aquatic Chesapeake Bay MD clone 3C002403 EU801
AF173821.1 gg_id:6994 Antarctic clone LB3-27 AF173
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EU636023.1 gg_id:277658 Antarctic str. CA1 EU636
EF520443.1 gg_id:230788 structure lakes: what controls ? acid-impacted lake clone ADK-BMe02-01 EF5204
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GQ030687.1 gg_id:456337 Topographical and Temporal Human Skin Microbiome skin palm clone nbw881e04c1 GQ030
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EF127603.1 gg_id:208255 Fossil Genes and Microbes Entombed Oldest Ice on Earth ice clone EME093 EF1276
CU927221.1 gg_id:549143 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU927
CT574057.1 gg_id:151642 Evry municipal wastewater treatment plant clone 054F01_B_DI_P58 CT574
CR933282.1 gg_id:109578 Evry municipal wastewater treatment plant clone 061G11_B_SD_P93 CR933
AF422679.1 gg_id:37583 TCE-sediment clone d156 AF4226
AY218554.1 gg_id:82194 penguin droppings sediments clone KD1-134 AY218
FJ982926.1 gg_id:422910 Simplicispira limi str. ST3 FJ9829
AF078755.1 gg_id:7088 Aquaspirillum psychrophilum str. LMG 5408 AF0787
AY780904.1 gg_id:143192 Aquaspirillum metamorphum str. 416 AY780
Z93960.1 gg_id:7097 activated sludge clone T33 Z93960
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EU652480.1 gg_id:278324 Simplicispira sp. str. JPB-3.01 EU6524
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CU466844.1 gg_id:324604 Evry wastewater treatment plant anoxic basin clone CU466
AJ863422.1 gg_id:126382 poplar tree microcosm rhizoplane flooded clone 23RPF1 AJ8634
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CU466904.1 gg_id:335159 Evry wastewater treatment plant anoxic basin clone CU466
CU466876.1 gg_id:316181 Evry wastewater treatment plant anoxic basin clone CU466
DQ640671.1 gg_id:233183 full-scale EBPR sludge clone Skagenf43 DQ640
AY275432.1 gg_id:98380 Brachymonas petroleovorans str. CHX AY2754
GQ024888.1 gg_id:484782 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ024

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FJ516827.1 gg_id:572684 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled lower s FJ5168
GQ102483.1 gg_id:433031 Topographical and Temporal Human Skin Microbiome skin axillary vault clone nbw7 GQ102
AF439400.1 gg_id:40752 Ramlibacter henchirensis str. TMB834 AF4394
AM411934.1 gg_id:237073 Variovorax sp. str. P-34 AM411
GU208252.1 gg_id:590127 Study on Sediment Dongping Lake sediment clone 6S3-1 GU208
AF144383.3 gg_id:7037 Ramlibacter tataouinensis str. TTB310 AF1443
FJ612171.1 gg_id:567473 and Biogeochemistry Dongping Lake Ecosystems lake water clone DP3.4.83 FJ6121
EF520494.1 gg_id:230979 structure lakes: what controls ? acid-impacted lake clone ADK-WYe02-39 EF5204
GQ022440.1 gg_id:487228 Topographical and Temporal Human Skin Microbiome skin axillary vault clone nbu31 GQ022
GQ020798.1 gg_id:488871 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ020
DQ264547.1 gg_id:141807 universal microarray broader than typical library when sampling subsurface groundw DQ264
EU978826.1 gg_id:544356 and metabolic potential metagenome glacier ice clone glb338c EU9788
FJ534997.1 gg_id:533886 Enhancement protein conversion volatile fatty acids accumulation during addition: ef FJ5349
GQ089363.1 gg_id:407227 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw3 GQ089
AJ565430.1 gg_id:89333 Hylemonella sp. WQH1 AJ5654
GQ059928.1 gg_id:403464 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ059
DQ234183.2 gg_id:142262 determined library mangrove clone DS100 DQ234
GQ060097.1 gg_id:404459 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ060
EF520474.1 gg_id:230959 structure lakes: what controls ? acid-impacted lake clone ADK-HDh02-57 EF5204
EU801607.1 gg_id:325206 It's all ranking aquatic Chesapeake Bay MD clone 3C002944 EU8016
FJ484871.1 gg_id:569745 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4848
GQ038137.1 gg_id:404029 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbu38 GQ038
EU801055.1 gg_id:320222 It's all ranking aquatic Delaware Bay NJ clone 2C229356 EU8010
GQ021512.1 gg_id:488156 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbu27 GQ021
EU881143.1 gg_id:323551 Karst region primary forest soil clone KPF200711-105 EU8811
EU652471.1 gg_id:279025 Pseudomonas sp. str. JPB-1.12 EU6524
AB166668.1 gg_id:101184 Leptothrix sp. str. SAYI-C AB1666
AF234683.1 gg_id:38002 sludge clone A15 AF2346
AJ867925.1 gg_id:143202 Fluctuations Significantly Influence Microbial Present Nival Lakes lake water clone D4 AJ8679
DQ640667.1 gg_id:229779 full-scale EBPR sludge clone Skagenf97 DQ640
GQ339130.1 gg_id:523295 Comprehensive Investigation on Iron Cycling Freshwater Seep ing Microscopy Cultiv GQ339
DQ521517.1 gg_id:190163 Antarctic lake ice cover clone ANTLV2_E04 DQ521
FM873335.1 gg_id:367349 occupant source house floor dust clone FD01H06 FM873
EF660749.1 gg_id:245337 Pelomonas soli str. SYB4 EF6607
EF632937.1 gg_id:252287 Unique microbial contrasting aquatic altitude Andean Altiplano (northern Chile) fresh EF6329
CU466870.1 gg_id:344265 Evry wastewater treatment plant anoxic basin clone CU4668
AB114618.1 gg_id:103415 rice plant clone ER-50 AB1146
EU542576.1 gg_id:258654 Ideonella sp. str. 1a22 EU5425
AF293003.1 gg_id:37495 ferromanganous micronodule clone MNE12 AF2930

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AJ422174.2 gg_id:23589 Germany:Spittelwasser River clone Spb283 AJ4221
DQ656489.1 gg_id:160616 Aquicola tertiaricarbonis str. L10 DQ656
FJ517015.1 gg_id:538250 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled water c FJ5170
EU881220.1 gg_id:355367 Karst region natural restoration land soil clone KNR200711-005 EU881
DQ413070.1 gg_id:152813 Microbial -dependent and Approaches Anaerobic/Aerobic SBR Reactor EBPR sludge DQ413
EF520457.1 gg_id:233054 structure lakes: what controls ? acid-impacted lake clone ADK-GRe02-95 EF5204
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AY947999.1 gg_id:143722 Synchrony and seasonality two temperate rivers USA: Massachusetts Parker River cl AY947
AY921693.1 gg_id:113507 farm soil clone AKYG1800 AY921
FJ612162.1 gg_id:576928 and Biogeochemistry Dongping Lake Ecosystems lake water clone DP3.4.31 FJ6121
EU638506.1 gg_id:340339 Ecological Role Firmicutes Identified Thermophilic Microbial Fuel Cells thermophilic n EU638
AB252904.1 gg_id:171114 Identification Shibayama lagoon Ishikawa Japan iron-oxidation biofilm clone 233 AB252
AY212560.1 gg_id:102288 water 10 m downstream manure clone 109ds10 AY212
EF540425.1 gg_id:218435 oil-shale semi-coke solid waste clone MK4a EF5404
FJ484912.1 gg_id:561277 microbial autonomous rover world's deepest phreatic sinkhole water column El Zacat FJ4849
DQ327690.1 gg_id:167798 Biological filtration limits carbon availability affects downstream biofilm formation a DQ327
DQ337559.1 gg_id:145288 Pseudomonas sp. BBCT8 DQ337
AM990747.1 gg_id:278347 Leptothrix sp. MOLA 523 AM990
FJ902318.1 gg_id:541652 phreatic limestone sinkholes northeastern Mexico biomat rock outcrop cenote La Pal FJ9023
FJ902628.1 gg_id:541372 phreatic limestone sinkholes northeastern Mexico biofilm 1m deep hydrothermal ver FJ9026
AY212645.1 gg_id:107717 water 20 m upstream manure clone 192up AY212
EU244077.1 gg_id:365878 16SrRNA probes river Leine sediment clone sls1359 EU244
AY947934.1 gg_id:143876 Synchrony and seasonality two temperate rivers USA: Massachusetts Ipswich River c AY947
AB240317.1 gg_id:144929 microbial structure Phragmites rhizosphere biofilm reed bed reactor laboratory clon AB240
AF234681.1 gg_id:48162 sludge clone A2 AF234
AY622242.1 gg_id:106999 subsurface soil clone B-E7 AY622
AB015048.1 gg_id:7128 Leptothrix sp. str. MBIC3364 Y8 AB015
DQ066964.1 gg_id:130439 Lake Washington sediment clone pLW-3 DQ066
DQ017912.1 gg_id:136963 upland stream clone S-Btb7_08 DQ017
DQ241397.1 gg_id:143057 Leptothrix sp. str. S1.1 DQ241
AB240488.1 gg_id:142529 microbial structure rhizosphere biofilm Sapporo root-tip (0 40 mm) Phragmites Sosei AB240
EU340201.1 gg_id:250274 on Aquatic Macrophytes Geothermally Active Area Sedge Bay Yellowstone Lake WY EU340
AF498703.1 gg_id:58127 soil isolate Ellin321 AF498
GQ093947.1 gg_id:404750 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ093

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AF234685.1 gg_id:17679 sludge clone A26b AF2346
FJ478604.1 gg_id:581639 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4786
EF079081.1 gg_id:244828 seeps sulfur-rich neutral pH ecosystem ochre-colored flocculent iron seep ditch fresh EF0790
AB240526.1 gg_id:144416 microbial structure rhizosphere biofilm Sapporo root-base (80 160 mm) Phragmites S AB2405
CU466896.1 gg_id:355908 Evry wastewater treatment plant anoxic basin clone CU4668
FJ152778.1 gg_id:341579 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX5A_ FJ1527
AM501430.1 gg_id:256189 Pelomonas saccharophila str. LMG2256 AM501
EU283393.1 gg_id:251129 consecutively aerated submerged treating domestic wastewater activated sludge m EU2833
AB480710.2 gg_id:556657 Leptothrix sp. str. PW5B AB4807
EU104010.1 gg_id:308805 Evidence Comamonadaceae determinants settling performance activated sludge clo EU1040
GQ402782.1 gg_id:557460 Accessing microbial biodiversity and metabolic potential tropical -based approach p GQ402
DQ836752.1 gg_id:169856 attached biomass denitrifying bioreactor clone DB-14 DQ836
CU466869.1 gg_id:333262 Evry wastewater treatment plant anoxic basin clone CU4668
X97071.1 gg_id:7126 Leptothrix mobilis str. Feox-1 X97071
AB271046.1 gg_id:172053 Leptothrix ginsengisoli str. Gsoil 1005 AB2710
AY212570.1 gg_id:107290 water 10 m downstream manure clone 119ds10 AY2125
CU466877.1 gg_id:305915 Evry wastewater treatment plant anoxic basin clone CU4668
EU015116.1 gg_id:243107 Investigation estrogen-degrading bioreactors membrane bioreactor suspension and EU0151
EU244040.1 gg_id:364166 16SrRNA probes river Leine sediment clone sl1288 EU2440
AY947983.1 gg_id:143699 Synchrony and seasonality two temperate rivers USA: Massachusetts Ipswich River c AY9479
EU340184.1 gg_id:235584 on Aquatic Macrophytes Geothermally Active Area Sedge Bay Yellowstone Lake WY EU3401
AJ576389.1 gg_id:104905 midgut homogenate Pachnoda ephippiata larva clone PeM24 AJ5763
EU817495.1 gg_id:334185 Comamonas sp. str. XJ-L79 EU8174
AY212575.1 gg_id:99071 water 10 m downstream manure clone 124ds10 AY2125
FJ823942.1 gg_id:529615 Controlling accumulation fermentation inhibitors biorefinery recycle water microbial FJ8239
EU801639.1 gg_id:342792 It's all ranking aquatic Chesapeake Bay MD clone 3C002983 EU8016
EU488023.1 gg_id:545997 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU4880
AM778004.1 gg_id:247071 nonsaline . Influence pore-size filters determination structure alkaline groundwater AM778
AY768824.1 gg_id:108787 deep terrestrial fracture system clone D8A_4 AY7688
DQ537534.1 gg_id:170339 manganese oxidation soil clone JH-GY34 DQ537
GQ263735.1 gg_id:558043 Structure Simulated Level Waste Site simulated level waste site clone WC1_a51 GQ263
DQ301787.1 gg_id:147885 Comamonas sp. str. PHD-10 DQ301
DQ824571.1 gg_id:191875 human fecal clone RL185_aaj69g09 DQ824
FJ009392.1 gg_id:305949 Hydrogenophaga intermedia str. C1 FJ0093
GQ404493.1 gg_id:588562 Hydrogenophaga flava str. ENV735 GQ404
AB195168.1 gg_id:108779 Comamonas sp. str. 147S AB1951
AF280845.1 gg_id:21592 bioreactor clone mle1-6 AF2808
AJ420328.1 gg_id:30654 Hydrogenophaga flava str. DSM 619 AJ4203
DQ202143.1 gg_id:139876 denitrifying treating groundwater contaminated nitrate and uranium fluidized bed r DQ202
EU037345.2 gg_id:261198 structure landfill sediments receiving long-term augmentations wastes chromium co EU0373

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DQ854970.1 gg_id:164474 Hydrogenophaga sp. str. GPTSA14 DQ854
AF019037.1 gg_id:6953 Hydrogenophaga intermedia str. S1 AF0190
AF508103.1 gg_id:66240 Variovorax paradoxus TG27 AF5081
DQ256326.1 gg_id:141917 subsurface water clone EV818BHEB5102502DRLWq27f021 DQ256
AM777949.1 gg_id:237044 nonsaline . Influence pore-size filters determination structure alkaline groundwater AM777
DQ824747.1 gg_id:173960 human fecal clone RL185_aaj72f07 DQ824
AB195164.1 gg_id:108969 Comamonas sp. str. PD-10t AB1951
AJ853614.1 gg_id:111232 landfill leachate clone GZKB122 AJ8536
AB294326.1 gg_id:204877 Microbial structure deep coal seem groundwater stream clone SWB15 AB2943
DQ234232.2 gg_id:141836 determined library mangrove clone DS150 DQ234
AY218568.1 gg_id:81130 penguin droppings sediments clone KD2-104 AY2185
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EF469609.1 gg_id:224686 Pseudacidovorax intermedius str. CC-CC21 EF4696
EF580985.1 gg_id:279450 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM5(EF5809
AY823974.1 gg_id:148639 Investigation Acetate-Fed Denitrifying Microbial Stable Isotope Probing Full-Cycle an AY8239
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AB077037.1 gg_id:78049 Macromonas bipunctata str. IAM 14880 AB0770
FJ382319.1 gg_id:572939 Potentially pathogenic and air stem cell transplant summer hospital shower water BM FJ3823
AY863093.1 gg_id:112155 wastewater treatment pool clone SXAU055 AY8630
DQ337093.1 gg_id:146820 subsurface water clone EV818SWSAP75 DQ337
DQ824754.1 gg_id:184303 human fecal clone RL185_aaj72g06 DQ824
AY947966.1 gg_id:141254 Synchrony and seasonality two temperate rivers USA: Massachusetts Ipswich River c AY9479
Z93993.1 gg_id:6940 activated sludge clone T60 Z93993
FJ598354.1 gg_id:552978 Study Prokaryotic Some Selected Niches Western Ghats India water clone BCWCWP1 FJ5983
AM110076.2 gg_id:250992 Hydrogenophaga sp. str. Rs71 AM110
AM042694.1 gg_id:161866 Tepidimonas thermarum str. AA2 AM042
AM777993.1 gg_id:251137 nonsaline . Influence pore-size filters determination structure alkaline groundwater AM777
AB271047.1 gg_id:171214 Hydrogenophaga sp. str. Gsoil 1545 AB2710
EF532793.1 gg_id:227420 Hydrogenophaga bisanensis str. K102 EF5327
FJ660572.1 gg_id:561367 Correlation functional stability and dynamics full-scale wastewater treatment plant a(FJ6605
GQ114408.1 gg_id:418439 Topographical and Temporal Human Skin Microbiome skin manubrium clone nbw59 GQ114
AF208386.1 gg_id:28592 Variovorax paradoxus str. WP1 AF2083
AY212671.1 gg_id:104415 water 20 m downstream manure clone 223ds20 AY2126
DQ824567.1 gg_id:191848 human fecal clone RL185_aaj69g05 DQ824
DQ349098.1 gg_id:148264 Variovorax limosa str. EMB320 DQ349
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AF509578.1 gg_id:91935 intestine clone AF5095

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AY622243.1 gg_id:107152 subsurface soil clone B-H11 AY622
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GQ115248.1 gg_id:465653 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ115
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EF157142.1 gg_id:254138 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF1571
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EF516381.1 gg_id:223622 grassland soil clone FCPS464 EF516
EF157145.1 gg_id:253142 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF157
EU104241.1 gg_id:351875 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
GQ002349.1 gg_id:507313 Topographical and Temporal Human Skin Microbiome skin buttock clone nbu176d0 GQ002
AB286592.1 gg_id:216359 Reducing production and domination comamonadaceae reducing oxygen supply was AB286
GQ060015.1 gg_id:448535 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ060
AY566579.1 gg_id:108727 Acidovorax sp. str. PG-06 AY566
GQ102770.1 gg_id:457672 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ102
EF492935.1 gg_id:255248 Iron-manganese Nodule and Its Surrounding Soil soil around iron-manganese nodule EF492
EU015090.1 gg_id:239385 Investigation estrogen-degrading bioreactors membrane bioreactor suspension and EU015
DQ451457.1 gg_id:156987 forest soil clone FAC18 DQ451
EU803951.1 gg_id:349195 It's all ranking aquatic Lake Gatun clone 5C231617 EU803
GQ039298.1 gg_id:450995 Topographical and Temporal Human Skin Microbiome skin axillary vault clone nbu41 GQ039
GQ008150.1 gg_id:501515 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ008
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GQ075567.1 gg_id:446083 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ075
FJ534986.1 gg_id:513449 Enhancement protein conversion volatile fatty acids accumulation during addition: ef FJ5349
GQ101831.1 gg_id:397741 Topographical and Temporal Human Skin Microbiome skin elbow clone nbw499g11 GQ101
CU918612.1 gg_id:539274 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU918
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DQ984534.1 gg_id:241827 FD oil-contaminated soil clone FD_17 DQ984
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GQ108883.1 gg_id:403824 Topographical and Temporal Human Skin Microbiome skin palm clone nbw718g05c GQ108
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EU800160.1 gg_id:353966 It's all ranking aquatic Delaware Bay NJ clone 2C228154 EU800
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DQ230947.1 gg_id:144759 subsurface water clone NO27FW100501SAB47 DQ230
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AY168733.2 gg_id:136160 Oxidizing Biofilms: Cultivation Approaches and Rates Arsenite Oxidation arsenite-oxi AY168
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GQ105741.1 gg_id:438927 Topographical and Temporal Human Skin Microbiome skin axillary vault clone nbw5 GQ105
GQ098862.1 gg_id:436788 Topographical and Temporal Human Skin Microbiome skin umbilicus clone nbw558a GQ098
EU676430.1 gg_id:302458 rhizosphere pioneer plants (Bahia xylopoda and Viguiera linearis) growing on metals EU676
GQ249361.1 gg_id:547824 Research on biodiversity sludge bioflim Biological Aeration Filter clone A17 GQ249
EU800119.1 gg_id:341614 It's all ranking aquatic Delaware Bay NJ clone 2C228107 EU800
EU801154.1 gg_id:314075 It's all ranking aquatic Delaware Bay NJ clone 2C229504 EU801
EU735706.1 gg_id:337175 Biological Arsenic Remediation Project Groundwater Aquifer soil clone 1As18 EU735
AF502223.1 gg_id:59344 EBPR sludge lab scale clone HP1B35 AF502
EF516428.1 gg_id:220686 grassland soil clone FCPP550 EF5164
GQ009720.1 gg_id:499945 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw77 GQ009
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DQ432053.1 gg_id:154227 Variovorax koreensis str. GH 9-3 DQ432
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AJ420329.1 gg_id:35360 Variovorax paradoxus str. DSM 66 AJ4203
EU169184.1 gg_id:247107 Variovorax paradoxus str. B54 EU169
DQ257419.1 gg_id:141979 Variovorax paradoxus str. CAI-26 DQ257
DQ256486.1 gg_id:144358 Variovorax paradoxus Jm111 DQ256
AB269775.1 gg_id:209527 Variovorax paradoxus str. HYK0512-SK12 AB269
EF641108.1 gg_id:279410 Variovorax paradoxus str. TBEA6 EF6411
FJ214196.1 gg_id:342494 Co-speciation earthworms and their nephridial symbionts nephridia clone Ai3-13 FJ2141
FJ214173.1 gg_id:327535 Co-speciation earthworms and their nephridial symbionts nephridia clone Lf3-5 FJ2141
FJ214186.1 gg_id:315987 Verminephrobacter sp. str. At4 FJ2141
AJ543434.1 gg_id:83899 nephridia Octolasion lacteum clone Ol2-2 AJ5434

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DQ327663.1 gg_id:146840 Verminephrobacter foetidai str. EF01-2 DQ327
DQ327662.1 gg_id:149224 Verminephrobacter foetidai str. EF01-1 DQ327
NC_008786.1 gg_id:179440 Verminephrobacter eiseniae str. EF01-2 NC_00
GQ066930.1 gg_id:374294 Topographical and Temporal Human Skin Microbiome skin back clone nbw730f08c1 GQ066
FJ605423.1 gg_id:430937 Xenophilus aerolata str. PR-1 FJ6054
NR_025114.1 gg_id:571390 Xenophilus azovorans str. KF46F; DSM 13620 NR_02
EF079087.1 gg_id:247480 Ideonella sp. str. K5.1 EF079
DQ664241.1 gg_id:164847 Ideonella sp. str. IMCC1722 DQ664
AB476705.1 gg_id:547214 SSU sequences prokaryotes inside emergency Gotthard road tunnel Switzerland sulf AB476
AJ421931.2 gg_id:51718 Germany:Elbe River clone Elb197 AJ4219
EU104142.1 gg_id:336502 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
EF572731.1 gg_id:231452 Coco's Island site 23 marine clone S23_830 EF5727
CU466908.1 gg_id:356048 Evry wastewater treatment plant anoxic basin clone CU466
EU340221.1 gg_id:251319 on Aquatic Macrophytes Geothermally Active Area Sedge Bay Yellowstone Lake WY EU340
NR_025916.1 gg_id:555018 Leptothrix discophora str. SS-1; ATCC 43182 NR_02
EU130970.1 gg_id:243909 Methylibium sp. BAC115 EU130
AM774413.1 gg_id:239835 Rivibacter subsaxonicus str. BF49 AM774
EU130974.1 gg_id:236153 Methylibium sp. BAC199 EU130
AJ422161.2 gg_id:25579 Germany:Spittelwasser River clone Spb111 AJ4221
EF516772.1 gg_id:225864 grassland soil clone FCPP576 EF5167
AM697195.1 gg_id:217417 sequencing libraries indoor dust clone BF0001C036 AM697
FM209333.1 gg_id:571602 Microbial Negev desert s sand soil clone 1250 FM209
AY382153.1 gg_id:136994 enriched nonylphenol ethoxylates activated sludge clone S1B10 AY382
AY218641.1 gg_id:81483 penguin droppings sediments clone KD4-64 AY2186
EU536904.1 gg_id:287179 profile human microbiota antecubital fossa (inner elbow) skin clone nbt223f06 EU536
AM777983.1 gg_id:239229 nonsaline . Influence pore-size filters determination structure alkaline groundwater AM777
FM209332.1 gg_id:537782 Microbial Negev desert s sand soil clone 225 FM209
EU499698.1 gg_id:274988 Field-based laboratory stable isotope probing surveys identities both aerobic and an EU499
EF515904.1 gg_id:226906 grassland soil clone FCPP759 EF5159
AY771860.1 gg_id:108580 wastewater treatment pool clone SXAU009 AY7718
DQ837237.1 gg_id:248403 Archaeal and pristine National Park Spain Donana coastal aquifer clone 49S1_2B_11 DQ837
AB240506.1 gg_id:142476 microbial structure rhizosphere biofilm Sapporo root-base (80 160 mm) Phragmites AB240
FJ893771.1 gg_id:540262 Matriptase-deficient mice exhibit ichthyotic selective shift skin microbiota B1E; ear pu FJ8937
AY921694.1 gg_id:114102 farm soil clone AKYH1130 AY9216
GU208509.1 gg_id:580636 Study on Sediment Dongping Lake sediment clone Shf4-24 GU208
AF280853.1 gg_id:24241 bioreactor clone mle1-19 AF2808
AY218657.1 gg_id:103477 penguin droppings sediments clone KD9-144 AY2186
EU907887.1 gg_id:309250 Identification hexahydro-135-trinitro-135-triazine (RDX)-degrading microorganisms v EU9078
EU362134.1 gg_id:305378 and Mesquite Flats Site Death Valley dune sand clone 5 EU362
EU491350.1 gg_id:259064 Abundance and microbial life ocean crust seafloor lavas Hawai'i South Point X3 clone EU491

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EU639232.1 gg_id:317724 Ecological Role Firmicutes Identified Thermophilic Microbial Fuel Cells thermophilic n EU6392
DQ298309.1 gg_id:153016 studies on and nifH Lasiurus sindicus Sewan grass Thar Desert Rajasthan India rhizos DQ298
FM872725.1 gg_id:366825 occupant source house floor dust clone FB01B01 FM872
EF515947.1 gg_id:223606 grassland soil clone FCPT445 EF5159
EF522266.1 gg_id:217402 Endolithic Microbial Ecosystems Rocky Mountain endolithic sandstone clone FQSS04 EF5222
EU881306.1 gg_id:304030 (maize-sweet cropping field) Karst region maize-sweet potato cropland soil clone KM EU8813
FJ478963.1 gg_id:511130 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4789
DQ640690.1 gg_id:234716 full-scale EBPR sludge clone Skagenf14 DQ640
EF516703.1 gg_id:226711 grassland soil clone FCPT716 EF5167
FJ592525.1 gg_id:547327 Fumarole-supported islands biodiversity within hyperarid -elevation landscape on Pu FJ5925
EF020021.1 gg_id:214611 aspen rhizosphere clone Elev_16S_1414 EF0200
EF580981.1 gg_id:278694 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM1 EF5809
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AY281150.1 gg_id:105543 Collimonas fungivorans str. NCCB 100029 AY2811
AJ496445.1 gg_id:57762 Collimonas fungivorans str. CTE227 = NCCB100027 AJ4964
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EU549858.1 gg_id:302270 Herbaspirillum frisingense str. AU14559 EU5498
EU399918.1 gg_id:275308 Herbaspirillum frisingense str. CAF265 EU3999
AB021424.1 gg_id:7295 Herbaspirillum rubrisubalbicans str. ATCC 19308T AB0214
AJ238356.1 gg_id:7299 Herbaspirillum rubrisubalbicans str. M4 AJ2383
AY191273.1 gg_id:80182 Herbaspirillum seropedicae str. Z152 AY1912

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AY191274.1 gg_id:80833 Herbaspirillum seropedicae str. ZA95 AY1912
AY191272.1 gg_id:80181 Herbaspirillum seropedicae str. ZA69 AY1912
AF361023.1 gg_id:74451 Herbaspirillum sp. str. NAH4 AF3610
AJ505863.1 gg_id:58574 Herbaspirillum sp. str. PIV-34-1 = DSM 14683 AJ5058
AY043371.1 gg_id:63615 Pseudomonas huttiensis str. AU2210A AY0433
GU433469.1 gg_id:528777 Herbaspirillum huttiense str. CCBAU 61429 GU433
EU024131.1 gg_id:243525 Herbaspirillum huttiense str. CIP 103296 EU0241
AJ583172.1 gg_id:94668 radioactive site ground water clone S15B-MN104 AJ5831
AY049039.1 gg_id:47876 Pseudomonas huttiensis str. D AY0490
AJ238359.1 gg_id:7302 Herbaspirillum frisingense str. Mb11 AJ2383
AB021366.1 gg_id:7292 Pseudomonas huttiensis str. ATCC 14670T AB0213
FJ266340.1 gg_id:534872 Herbaspirillum rubrisubalbicans str. RFNB30 FJ2663
FJ605419.1 gg_id:404943 Herbaspirillum rubrisubalbicans str. PPs-3 FJ6054
NR_028656.1 gg_id:584170 Herbaspirillum putei str. 7-2; IAM 15032 NR_028
AY191276.2 gg_id:80103 Herbaspirillum seropedicae str. M2 AY1912
AB512142.1 gg_id:564904 Herminiimonas fonticola str. da Costa S-94 (=CIP 108398 =LMG AB5121
AB512141.1 gg_id:546345 Herminiimonas saxobsidens str. NEU 1218 (=DSM 18748 =CCM 7436) AB5121
EU133873.1 gg_id:251868 tallgrass prarie soil clone FFCH17907 EU1338
GU244359.1 gg_id:593835 Janthinobacterium sp. str. Lc10-8 GU244
EU847234.1 gg_id:353532 Janthinobacterium sp. str. blxAG EU8472
FJ894020.1 gg_id:555629 Matriptase-deficient mice exhibit ichthyotic selective shift skin microbiota B1E; ear pu FJ8940
FJ193906.1 gg_id:313175 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1939
EU536428.1 gg_id:285908 profile human microbiota antecubital fossa (inner elbow) skin clone nbt213f12 EU5364
AF529096.1 gg_id:62419 trichloroethene-contaminated site clone ftl206 AF5290
EU982453.1 gg_id:341939 Enrichment and detection moderate thermophilic microorganisms vents Lau Basin h EU9824
AY792243.1 gg_id:119507 Humic Lake clone CrystalBog5B11 AY7922
FJ267554.1 gg_id:308624 Massilia sp. str. II_Gauze_W_12_10 FJ2675
DQ228653.1 gg_id:142568 Microbial colonizing mineral surfaces within sulfide-microbial incubator clone CH2b1 DQ228
FM209304.1 gg_id:571521 Microbial Negev desert s sand soil clone 2321 FM209
AM412135.1 gg_id:241097 Massilia sp. str. P-118 AM412
GQ354568.1 gg_id:556334 Naxibacter sp. str. HMD2167 GQ354
EU221406.1 gg_id:235543 Massilia timonae str. H2P8 EU2214
GQ015305.1 gg_id:494362 Topographical and Temporal Human Skin Microbiome skin palm clone nbw766g11c1 GQ015
FM209303.1 gg_id:536558 Microbial Negev desert s sand soil clone 2220 FM209
GQ098872.1 gg_id:406246 Topographical and Temporal Human Skin Microbiome skin umbilicus clone nbw558b GQ098
AB487721.1 gg_id:539718 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB4877
AY792245.1 gg_id:126195 Humic Lake clone CrystalBog2G3 AY7922
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EU244075.1 gg_id:365805 16SrRNA probes river Leine sediment clone sl1354 EU2440

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EU249991.1 gg_id:257841 Massilia sp. str. PmeaMuc24 EU2499
FJ191812.1 gg_id:323439 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1918
AB488284.1 gg_id:569272 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR0 AB4882
EU626407.1 gg_id:279407 Massilia sp. str. biosZ4 EU6264
EU373360.1 gg_id:270893 Massilia timonae str. HNL19 EU3733
FJ605429.1 gg_id:456558 Janthinobacterium sp. str. PR-2 FJ6054
FJ191801.1 gg_id:331807 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1918
EU365682.1 gg_id:269884 Massilia timonae str. M-10 EU3656
EU160504.1 gg_id:318413 diagnosis central-venous infections cultivation-dependent and techniques biofilm sc EU1605
EF682966.1 gg_id:246159 Eastern Mediterranean atmosphere clone F27_2E_FL EF6829
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AB487246.1 gg_id:558053 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNIR0 AB4872
FJ375473.1 gg_id:510182 T-RFLP powerful tool rapid assessment cells microbial fuel cell sheet carbon paper an FJ3754
AB234525.1 gg_id:145411 termite gut homogenate clone MgMjD-103 AB2345
EU274637.2 gg_id:259460 Massilia timonae str. U2 EU2746
AY966001.1 gg_id:115346 Sinobacter luteus str. 101 AY9660
AB487486.1 gg_id:567594 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB4874
EU431713.1 gg_id:540535 Troglobitic (Cansiliella) on specialized feeding apparatus calcium carbonate (moonmi EU4317
EF516951.1 gg_id:223889 grassland soil clone FCPU439 EF5169
AJ871463.1 gg_id:110987 Massilia timonae CP183-9 AJ8714
GU377105.1 gg_id:518781 Janthinobacterium sp. str. sptzw14 GU377
AF443584.1 gg_id:88301 semiarid soil clone C-F-25 AF4435
AM412137.1 gg_id:236550 Massilia sp. str. P-127 AM412
AF408326.1 gg_id:34181 Massilia sp. clone KL-11-1-5 AF4083
AM412141.1 gg_id:235451 Oxalobacter sp. str. P-154 AM412
AY512824.1 gg_id:106310 Massilia timonae str. S11 AY5128
EU554441.1 gg_id:275614 Naxibacter haematophilus str. Ag15 EU5544
AY679161.1 gg_id:102382 Naxibacter alkalitolerans str. YIM 31775 AY6791
AB097172.1 gg_id:87579 Janthinobacterium sp. str. J3 AB0971
EU369116.1 gg_id:551576 Janthinobacterium sp. str. AR12 EU3691
AY599723.1 gg_id:102086 fungal ascocarp clone Cl-73-TB2-II AY5997
EU861944.1 gg_id:344291 effects chronic fertilization on alpine tundra microbial implications carbon and cyclin EU8619
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EU431682.1 gg_id:557117 Troglobitic (Cansiliella) on specialized feeding apparatus calcium carbonate (moonmi EU4316
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GQ339887.1 gg_id:580212 Massilia sp. str. MK02 GQ339
U49752.1 gg_id:70632 Oxalobacter formigenes str. OXWR wild rat cecum U49752
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U49749.2 gg_id:7365 Oxalobacter formigenes str. BA4 human feces U4974
U49758.1 gg_id:7372 Oxalobacter formigenes str. HC1 human feces U4975
U49756.1 gg_id:7371 Oxalobacter formigenes str. OXK human feces U4975
FJ264587.1 gg_id:331690 Manganese and Iron Dependent Anaerobic Oxidation Methane methane seep sedim FJ2645
AB488167.1 gg_id:559492 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR AB488
U49754.1 gg_id:7369 Oxalobacter formigenes str. OXCR laboratory rat cecum U4975
AB486719.1 gg_id:591499 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSCOR AB486
FJ374254.1 gg_id:363874 Inter- and intraspecific humivorous scarab beetle larvae (Pachnoda spp.) larval hindg FJ3742
EF454731.2 gg_id:217779 termite hindgut clone 290cost002-P3L-2608 EF4547
EF434363.1 gg_id:216215 human infant gastrointestinal tract microbiota stool mother babies 13 and 14 7 mont EF4343
AB288876.1 gg_id:202850 termite gut homogenate clone BOf1-03 AB288
DQ796767.1 gg_id:174649 human fecal clone RL179_aao54b12 DQ796
FJ269064.1 gg_id:340557 dissimilatory ferric iron reducers arsenic contaminated Hunan south China paddy soil FJ2690
DQ676429.1 gg_id:224948 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676
AB038368.1 gg_id:19906 Diaphorina citri symbiont AB038
FJ371291.1 gg_id:360508 feces adult twins and mothers; TS6 clone TS6_a03c02 FJ3712
AB234524.1 gg_id:145512 termite gut homogenate clone MgMjD-092 AB234
AF371862.1 gg_id:50320 swine intestine clone p-2518-18B5 AF371
X92555.1 gg_id:7343 Paucimonas lemoignei str. LMG 2207 T X9255
AB487570.1 gg_id:592988 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR AB487
GQ022806.1 gg_id:486864 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu3 GQ022
AY177773.1 gg_id:78721 soil clone AY177
GQ069768.1 gg_id:401837 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ069
DQ248254.1 gg_id:141616 carbon tetrachloride contaminated soil clone UE7 DQ248
AY509446.2 gg_id:99154 freshwater clone LiUU-5-340 AY509
DQ351915.1 gg_id:172665 manganese oxidation soil clone JH-WH04 DQ351
GU213338.1 gg_id:593816 Janthinobacterium sp. str. 338 GU213
AM292626.1 gg_id:173162 Influence uranyl and sodium nitrate on natural soil uranium mining waste pile clone AM292
DQ537535.1 gg_id:162048 manganese oxidation soil clone JH-GY37 DQ537
AY167838.1 gg_id:101564 Janthinobacterium agaricidamnosum str. SAFR-022 AY167
AB240320.1 gg_id:144767 microbial structure Phragmites rhizosphere biofilm reed bed reactor laboratory clon AB240
AB487490.1 gg_id:569658 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR AB487
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AM888196.1 gg_id:247545 Herbaspirillum sp. str. THWCSN35 AM888
EF516890.1 gg_id:226310 grassland soil clone FCPU733 EF5168
DQ906892.1 gg_id:256121 Oman subsurface soil clone 12A-1 DQ906
AY080915.1 gg_id:61192 acidic forest soil clone UP9 AY0809
AM991238.1 gg_id:279160 Microbial groundwater and their potential use biomonitoring karst spring water clo AM991
AF468345.1 gg_id:79836 Arctic sea ice ARK10157 AF4683

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EF516112.1 gg_id:225949 grassland soil clone FCPT483 EF5161
FM874576.1 gg_id:357346 occupant source house mattress dust clone MD03G10 FM874
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EF516978.1 gg_id:222289 grassland soil clone FCPU492 EF5169
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FM209308.1 gg_id:566578 Microbial Negev desert s sand soil clone 3303 FM209
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GU295961.1 gg_id:585352 Oxalicibacterium sp. str. JCN-21 GU295
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DQ787695.1 gg_id:166885 holocene lake sediment clone SLB419 DQ787
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FM872500.1 gg_id:362441 occupant source house floor dust clone FA01C01 FM872
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AB487777.1 gg_id:579702 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB487
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DQ490985.1 gg_id:157892 Pseudoburkholderia malthae str. CC-AFH3 DQ490
FM955855.1 gg_id:590236 Massilia sp. str. Asd M1A2 FM955
EU735672.1 gg_id:345327 archaeal structures oil contaminated and soils China pristine soil Jidong Oilfield clone EU735
EU636042.1 gg_id:279632 Antarctic str. N14 EU636
EU362137.1 gg_id:352209 and Mesquite Flats Site Death Valley dune sand clone 8 EU362
AY198108.1 gg_id:80296 melt pond on Arctic sea ice floe clone ARKMP92 AY198
FJ191604.1 gg_id:340957 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1916
AY367012.1 gg_id:93490 ultramicrobacterium str. Um1 AY367
EF682968.1 gg_id:255137 Eastern Mediterranean atmosphere clone F13_2C_FL EF6829
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GU124500.2 gg_id:570154 endosymbiotic (Stal) (rice brown planthopper) Endosymbiont Nilaparvata lugens clo GU124
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FM253595.1 gg_id:341548 Rock gold mine unusually microbial biofilm clone D06-1 FM253
AB488125.1 gg_id:548180 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR0 AB488
GQ098449.1 gg_id:401707 Topographical and Temporal Human Skin Microbiome skin umbilicus clone nbw559g GQ098
GQ461666.1 gg_id:533088 Coupling -ITS Clone Libraries and Automated Intergenic Spacer Microbial Bioremedi GQ461
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DQ248255.1 gg_id:142344 carbon tetrachloride contaminated soil clone UA6 DQ248
AF358019.1 gg_id:26437 peat soil microcosm clone LO13.5 AF358
DQ469209.1 gg_id:157564 Microbial and ecology water Zhong Tiaoshan mine Shanxi province China AMD clone DQ469
AF529095.1 gg_id:62422 trichloroethene-contaminated site clone FTL205 AF529
EF620474.1 gg_id:248260 ultramicrobacterium str. ISSDS-831 EF6204
NR_028859.1 gg_id:566047 Herbaspirillum lusitanum str. P6-12; LMG 21710 NR_02
DQ017931.1 gg_id:133568 upland stream clone S-Jos_62 DQ017
DQ787707.1 gg_id:171969 holocene lake sediment clone SLB523 DQ787
EU676429.1 gg_id:287560 rhizosphere pioneer plants (Bahia xylopoda and Viguiera linearis) growing on metals- EU676
EF592558.1 gg_id:232495 Duganella sp. str. B2 EF5925
AJ582194.1 gg_id:94238 soil near uranium mill tailings clone Gitt-GS-138 AJ5821
DQ177470.1 gg_id:140398 Janthinobacterium sp. str. Tibet-IIK43 DQ177
EU528198.1 gg_id:286535 Seasonal Dynamics Mudflat Mouth Major Kentucky Lake Reservoir Tributary sedime EU528
DQ341421.1 gg_id:150786 Janthinobacterium sp. str. Antarctic IS09 DQ341
AB008503.1 gg_id:7288 ultramicrobacterium str. MY14 AB008
EF682967.1 gg_id:243222 Eastern Mediterranean atmosphere clone F14_4D_FL EF6829
EU669620.1 gg_id:329777 soil clone S22-8 EU669
DQ158104.1 gg_id:139019 Use Field-Based Stable Isotope Probing Identify Adapted and Track Carbon F throug DQ158
EU636048.1 gg_id:278716 Antarctic str. GAo51 EU636
EF019643.1 gg_id:215915 aspen rhizosphere clone Elev_16S_794 EF0196
EU489741.1 gg_id:277881 Herminiimonas sp. str. UMB49 EU489
EU636041.1 gg_id:278413 Antarctic str. GA091 EU636
AF407411.1 gg_id:21324 MCB-contaminated groundwater-treating reactor clone RB13C5 AF4074
AY214204.1 gg_id:81252 benzene-contaminated groundwater clone ZZ15C3 AY2142
EU057874.1 gg_id:253699 Duganella sp. 5B EU057
AB488131.1 gg_id:589716 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR AB488
EF515920.1 gg_id:218687 grassland soil clone FCPT636 EF5159
AY218774.1 gg_id:81640 penguin droppings sediments clone KD6-99 AY2187
DQ188985.1 gg_id:140045 Herbaspirillum rhizosphaerae str. UMS-40 DQ188
FJ391493.1 gg_id:364438 primers targeting Gallionella- neutrophilic oxidizers enrichments and soils iron rich irr FJ3914
GU205377.1 gg_id:583924 Massilia sp. str. DR-a8 GU205
AY214210.1 gg_id:81997 benzene-contaminated groundwater clone ZZ17C9 AY2142
AF468324.1 gg_id:80693 melt pond on Arctic sea ice floe clone ARKMP-20 AF4683
AJ867677.1 gg_id:109615 melted red snow water clone RS 8-Bact69 AJ8676
AJ295498.1 gg_id:27239 rape rhizosphere clone wr0040 AJ2954
EU194890.1 gg_id:359091 Duganella sp. str. B41 EU1948
AM085762.1 gg_id:138928 Herminiimonas aquatilis str. CCUG 36956 AM085
GU377114.1 gg_id:525330 Janthinobacterium sp. str. sptzw26 GU377
GQ451298.1 gg_id:585579 Fecal Yunnan snub-nosed monkey feces clone J308 GQ451
AB487141.1 gg_id:588705 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNIR AB487

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AB487627.1 gg_id:544493 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB4876
AB487567.1 gg_id:588968 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB4875
GU213421.1 gg_id:589183 Janthinobacterium sp. str. 286 GU213
EU150263.1 gg_id:278184 acidobacteria and soil Niwot Ridge LTER clone Ni_E6P2 EU150
AY198109.1 gg_id:79646 melt pond on Arctic sea ice floe clone ARKMP91 AY1981
EF018240.1 gg_id:207516 aspen rhizosphere clone Amb_16S_591 EF0182
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AF468329.1 gg_id:80338 melt pond on Arctic sea ice floe clone ARKMP-56 AF4683
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EU636047.1 gg_id:278100 Antarctic str. GA045 EU636
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DQ264563.1 gg_id:141833 universal microarray broader than typical library when sampling subsurface groundv DQ264
EF574525.1 gg_id:232833 site S25 near Coco's Island marine clone S25_869 EF5745
AB487774.1 gg_id:583648 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB4877
AY061962.2 gg_id:202769 Oxalicibacterium flavum str. TA17; NEU98; LMG21571 AY0619
EU881344.1 gg_id:329302 (maize-sweet cropping field) Karst region maize-sweet potato cropland soil clone KM EU8813
AB488296.1 gg_id:587787 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR0 AB4882
DQ787694.1 gg_id:162906 holocene lake sediment clone SLB418 DQ787
EU160577.1 gg_id:310584 diagnosis central-venous infections cultivation-dependent and techniques biofilm sc EU1605
AB488003.1 gg_id:580561 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR0 AB4880
AM697274.1 gg_id:222450 sequencing libraries indoor dust clone BF0001C115 AM697
AB074523.1 gg_id:66301 Aquaspirillum arcticum str. IAM 14963 AB0745
AJ867657.1 gg_id:110953 melted red snow water clone RS 8-Bact21 AJ8676
AJ532689.1 gg_id:76765 uranium mill tailings clone Gitt-KF-179 AJ5326
GQ080374.1 gg_id:373791 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ080
AB074524.1 gg_id:56608 Aquaspirillum autotrophicum str. IAM 14942 AB0745
AY792246.1 gg_id:122054 Humic Lake clone CrystalBog5F8 AY7922
FJ517670.1 gg_id:568152 Disturbing epithelial homeostasis metazoan Hydra leads drastic changes microbiota e FJ5176
GU213418.1 gg_id:572788 Janthinobacterium sp. str. 278 GU213
AB487887.1 gg_id:587586 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR0 AB4878
GU113050.1 gg_id:583060 soil overlying mountain Manasi County Xinjiang soils influenced underground coal m GU113
AY515386.1 gg_id:111167 sediment Lake Tuusulanjarvi isolate str. F39 AY5153
AM849431.1 gg_id:236633 Seasonality Piburger See clone PIB-20 AM849
FJ535228.1 gg_id:572527 Contamination risk vegetables Arcobacter during post-harvest processing water carr FJ5352
AY227003.1 gg_id:81741 melt pool on Arctic sea ice floe clone ARKMP-94 proteobacterium AY2270
EF094990.1 gg_id:194580 Rubrivivax gelatinosus str. L31 EF0949
CU466884.1 gg_id:313285 Evry wastewater treatment plant anoxic basin clone CU466

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FJ484872.1 gg_id:512841 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4848
GQ396839.1 gg_id:587796 Functional unvegetated perhumid recently-deglaciated soils do not correlate shifts s GQ396
CU466928.1 gg_id:325979 Evry wastewater treatment plant anoxic basin clone CU466
GU454909.1 gg_id:528368 Understanding short-chain fatty acids accumulation enhanced alkaline digestion: kir GU454
CU466864.1 gg_id:327001 Evry wastewater treatment plant anoxic basin clone CU466
CU466850.1 gg_id:338411 Evry wastewater treatment plant anoxic basin clone CU466
GQ249369.1 gg_id:553551 Research on biodiversity sludge bioflim Biological Aeration Filter clone B11 GQ249
GQ302537.1 gg_id:585468 prokaryotic sediment Shawan methods cold spring clone sw-xj209 GQ302
AB087565.1 gg_id:80525 Sphaerotilus sp. L8 AB087
AB087561.1 gg_id:80068 Sphaerotilus sp. L2 AB087
Z18534.1 gg_id:7139 Sphaerotilus natans str. 565 Z18534
AF309815.1 gg_id:73442 synthetic wastewater tmbr15-22 complete clone wastewater AF3098
AF177943.1 gg_id:42541 Tepidimonas ignava str. SPS-1037T AF177
EF648100.1 gg_id:255518 dynamics during produced water treatment aerobic activated sludge clone HB116 EF6481
AY845054.1 gg_id:110733 Tepidimonas sp. str. I1-1 AY845
AY594193.1 gg_id:124693 Tepidimonas arfidensis AY594
EU799903.1 gg_id:338160 It's all ranking aquatic Newport Harbour RI clone 1C227573 EU799
AY519354.1 gg_id:107473 Thiomonas sp. str. 51 AY519
FM992405.1 gg_id:536889 Thiomonas intermedia FM992
AY455808.1 gg_id:103970 Thiomonas perometabolis str. ATCC 23370 AY455
DQ146028.1 gg_id:138509 Thiomonas sp. ML1-60 DQ146
DQ146101.1 gg_id:138673 Thiomonas sp. ML2-48 DQ146
DQ146068.1 gg_id:139031 Thiomonas sp. ML2-1 DQ146
AB462557.1 gg_id:321818 Microbial production and energy source hyperthermophilic prokaryotes geothermal AB462
DQ146030.1 gg_id:138725 Thiomonas sp. ML1-62 DQ146
DQ146119.1 gg_id:138822 Thiomonas sp. ML2-67 DQ146
DQ146041.1 gg_id:138202 Thiomonas sp. ML1-74 DQ146
AB308270.1 gg_id:253106 Thiomonas sp. str. RAN4 AB308
DQ146097.1 gg_id:138841 Thiomonas sp. ML2-42 DQ146
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DQ902578.1 gg_id:189483 Thiomonas sp. str. CS2 DQ902
DQ146103.1 gg_id:138853 Thiomonas sp. ML2-5 DQ146
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DQ146130.1 gg_id:138505 Thiomonas sp. ML2-77 DQ146
DQ146065.1 gg_id:138239 Thiomonas sp. ML1-97 DQ146
DQ092334.1 gg_id:181396 Thiomonas sp. str. S10 DQ092

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GU233443.1 gg_id:531748 Thiomonas thermosulfata str. 21071 32621 GU233
EU937876.1 gg_id:351331 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU937
AY050601.1 gg_id:53885 MCB-contaminated groundwater-treating reactor clone GOUTB18 AY050
GU208450.1 gg_id:536615 Study on Sediment Dongping Lake sediment clone Fr2-13 GU208
EU937989.1 gg_id:307219 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU937
EF520449.1 gg_id:229543 structure lakes: what controls ? acid-impacted lake clone ADK-BTh02-80 EF5204
GQ388859.1 gg_id:562874 quality deterioration drinking system during serious red water outbreak clone Z238 GQ388
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FJ391492.1 gg_id:367650 primers targeting Gallionella- neutrophilic oxidizers enrichments and soils iron rich irr FJ3914
AB475012.1 gg_id:540849 sequences located Engiadina Bassa Valley Switzerland iron containing freshwater so AB475
EU937953.1 gg_id:356589 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU937
FM204958.1 gg_id:576266 Prokaryote shifts and reductive dehalogenase redundancy response electron donor FM204
GU208476.1 gg_id:592888 Study on Sediment Dongping Lake sediment clone Shc3-16 GU208
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GQ388776.1 gg_id:576649 quality deterioration drinking system during serious red water outbreak clone Z10 GQ388
EU937859.1 gg_id:327025 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU937
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GQ388790.1 gg_id:593879 quality deterioration drinking system during serious red water outbreak clone Z114 GQ388
AY766002.1 gg_id:108663 streamer acidic iron-rich spa water clone TrefC4 AY766
FJ037638.1 gg_id:310147 biofilms growing tropical warm soda spring Grenada West Indies clone 070218-Grena FJ0376
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AM167968.1 gg_id:143643 flora waters upstreaming fluids deep igneous rock aquifers mineral spring clone BB2 AM167
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EF562088.1 gg_id:241212 Subsurface microbial deep-granitic-fracture water colorado Henderson Molybdenum EF562
GQ339166.1 gg_id:520139 Comprehensive Investigation on Iron Cycling Freshwater Seep ing Microscopy Cultiv GQ339
EU283473.1 gg_id:278266 Model Early Earth Harbors Deeply Divergent Divisions Anderson Lake clone A531 EU283
GU183583.1 gg_id:551555 Structures Reduced and Oxidized Aquifer lower Aqueous Arsenic Titas Bangladesh a GU183
DQ676395.1 gg_id:225924 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676

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EF589997.2 gg_id:234236 changes structure sediments Songhua River contaminated nitrobenzene -DGGE cloni EF5899
EU937884.1 gg_id:325090 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU9378
GQ263781.1 gg_id:541843 Structure Simulated Level Waste Site simulated level waste site clone WC1_b19 GQ263
DQ676316.1 gg_id:223362 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676
GQ264072.1 gg_id:593060 Structure Simulated Level Waste Site simulated level waste site clone WC3_170 GQ264
FN430666.1 gg_id:540799 and activity autotrophic Fe(II)-oxidizing nitrate-reducing enrichment clone F29F67 FN4306
EU937881.1 gg_id:336779 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU9378
DQ386264.1 gg_id:196465 Siderooxidans lithoautotrophicus str. ES-1 DQ386
GQ388881.1 gg_id:579184 quality deterioration drinking system during serious red water outbreak clone Z40 GQ388
EU937923.1 gg_id:336177 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU9379
GQ389039.1 gg_id:553899 quality deterioration drinking system during serious red water outbreak clone S101 GQ389
EU937931.1 gg_id:339523 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU9379
DQ640694.1 gg_id:228790 full-scale EBPR sludge clone Skagenf48 DQ640
EU030485.1 gg_id:242960 microbial beneath sheet west Antarctic ice sheet; KIS sediment clone KIS.B77 EU0304
AY662039.1 gg_id:137480 uranium-contaminated groundwater clone 015B-E09 AY6620
AF351237.1 gg_id:16839 obtained coal tar waste-contaminated groundwater clone 8-11 AF3512
AB176698.1 gg_id:98919 subsurface geothermal water clone HAuD-LB/2-20 AB1766
DQ386858.1 gg_id:178053 Siderooxidans paludicola str. BrT DQ386
AY050584.1 gg_id:21368 MCB-contaminated groundwater-treating reactor clone GOUTA12 AY0505
GQ389053.1 gg_id:563337 quality deterioration drinking system during serious red water outbreak clone S114 GQ389
FN430662.1 gg_id:556677 and activity autotrophic Fe(II)-oxidizing nitrate-reducing enrichment clone F25F63 FN4306
AB252929.1 gg_id:164438 Identification Shibayama lagoon Ishikawa Japan iron-oxidation biofilm clone 472 AB2529
AB240306.1 gg_id:142150 microbial structure Phragmites rhizosphere biofilm reed bed reactor laboratory clone AB2403
FJ037639.1 gg_id:339910 biofilms growing tropical warm soda spring Grenada West Indies clone 070623-Grena FJ0376
GU220769.1 gg_id:535045 Microbial mineralization and involved microbial Basin field temperature hydrotherm GU220
GU183582.1 gg_id:573953 Structures Reduced and Oxidized Aquifer lower Aqueous Arsenic Titas Bangladesh a GU183
AY532538.1 gg_id:104211 uranium-contaminated aquifer clone 1013-1-CG10 AY5325
GQ389139.1 gg_id:512437 quality deterioration drinking system during serious red water outbreak clone S202 GQ389
DQ450774.1 gg_id:159803 Microbial Alpine Tundra Wet Meadow Soil: Chloroflexi Cold Water-Saturated satura DQ450
EF368015.1 gg_id:214563 Hydrogenophilus halorhabdus str. GI-445 EF3680
EF368017.1 gg_id:205395 Hydrogenophilus halorhabdus str. 1438 EF3680
NR_024663.1 gg_id:575778 Hydrogenophilus thermoluteolus str. TH-1; IFO 14978 NR_024
AY222297.1 gg_id:85885 silica sinter depositing geothermal power station discharge drain clone ST01-SN1D pro AY2222
AY820184.1 gg_id:109148 Hydrogenophilus denitrificans str. BDN3 AY8201
AF402975.1 gg_id:40122 hot spring clone bacteriap16 AF4029
NC_007404.1 gg_id:138944 Thiobacillus denitrificans str. ATCC 25259 NC_007
GU208337.1 gg_id:576708 Study on Sediment Dongping Lake sediment clone Se1-13 GU208
GQ472408.1 gg_id:555271 Microbial Dongping Lake lake sediment clone 4Db1-18 GQ472
AM936009.1 gg_id:572819 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM936
AB476681.1 gg_id:563379 SSU sequences prokaryotes inside emergency Gotthard road tunnel Switzerland sulf AB4766

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FJ167506.1 gg_id:348470 microbiology granular sludge under autotrophic mixotrophic and heterotrophic denit FJ1675
EU662520.1 gg_id:589721 Productivity- Chemolithoautotrophically Based Sulfidic Karst Systems floating microb EU662
GQ472333.1 gg_id:542530 Microbial Dongping Lake lake sediment clone 2S4-9 GQ472
FM213044.1 gg_id:568820 Biodiversity and shifts assessment biology tools Biotrickling treating loads biotricklin FM213
DQ390445.1 gg_id:154067 Thiobacillus sayanicus str. 4HG DQ390
DQ676323.1 gg_id:225078 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676
DQ499276.1 gg_id:159653 acidic cave snottite biofilm clone CV10 DQ499
EU735730.1 gg_id:354395 Biological Arsenic Remediation Project Groundwater Aquifer soil clone C_G46 EU735
EU735731.1 gg_id:351349 Biological Arsenic Remediation Project Groundwater Aquifer soil clone C_H29 EU735
DQ088733.1 gg_id:133752 subsurface clone BE16FW031601GDW_hole1-31 DQ088
AY955090.1 gg_id:114460 sediment clone 35-16K AY955
AJ243144.1 gg_id:23915 Thiobacillus denitrificans str. NCIMB 9548 AJ2431
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AB179662.1 gg_id:104873 groundwater 0.2 m-filterable fraction clone KNA6-NB06 AB179
AM934951.1 gg_id:540782 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM934
GU208503.1 gg_id:584889 Study on Sediment Dongping Lake sediment clone Shf3-3 GU208
AB179683.1 gg_id:106165 groundwater 0.2 m-captured fraction clone KNA6-EB03 AB179
FM213007.1 gg_id:570827 Biodiversity and shifts assessment biology tools Biotrickling treating loads biotricklin FM213
DQ415787.1 gg_id:157109 Dominant microbial limestone-corroding biofilms frasassi system Italy Frasassi sulfid DQ415
AY133108.1 gg_id:59893 TCE-contaminated site clone ccspost271 AY133
GU208385.1 gg_id:583159 Study on Sediment Dongping Lake sediment clone Td2-18 GU208
EU676425.1 gg_id:295407 rhizosphere pioneer plants (Bahia xylopoda and Viguiera linearis) growing on metals- EU676
GU208247.1 gg_id:549869 Study on Sediment Dongping Lake sediment clone 6S1-11 GU208
GU208366.1 gg_id:537091 Study on Sediment Dongping Lake sediment clone Td1-5 GU208
AY532567.1 gg_id:101367 uranium-contaminated aquifer clone 1013-28-CG14 AY532
DQ228387.1 gg_id:142824 microbial compositions two subglacial possible role microbes chemical weathering p DQ228
AB181501.1 gg_id:109267 wastewater biofilm clone SN-84 AB181
DQ988290.1 gg_id:180796 Microbial lab-scale A1-A2-O fixed biofilm system coking wastewater treatment with DQ988
FJ982929.1 gg_id:406533 Thiobacillus sp. str. ST22 FJ9829
GU208288.1 gg_id:576433 Study on Sediment Dongping Lake sediment clone Ftb3-6 GU208
GQ472327.1 gg_id:571504 Microbial Dongping Lake lake sediment clone 2S3-14 GQ472
GQ921437.1 gg_id:578101 Worms hell: presence Nematoda terrestrial deep subsurface (0.9-3.6 km) fracture w GQ921
DQ415784.1 gg_id:158397 Dominant microbial limestone-corroding biofilms frasassi system Italy Frasassi sulfid DQ415
DQ256331.1 gg_id:141880 subsurface water clone EV818BHEB5102502DRLWq27f055 DQ256
DQ088746.1 gg_id:129993 subsurface clone BE23FW031301A4RD-12 DQ088
GQ487949.1 gg_id:593089 Microbial metal soil polluted heavy metals clone T7-107 GQ487
AJ307935.1 gg_id:18168 derived microbial 'pearl'- community clone sipK89 AJ3079
AM934969.1 gg_id:535954 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM934
AB354624.1 gg_id:260306 Laminated iron texture iron-oxidizing calcite travertine iron-rich deposit clone FD015 AB354
EF079668.1 gg_id:252105 Thiobacillus sp. K6.2 EF0796

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AJ295643.1 gg_id:17101 uranium mill tailing clone KF-Gitt2-40 AJ2956
GU208480.1 gg_id:576640 Study on Sediment Dongping Lake sediment clone Shc1-6 GU208
AB425057.1 gg_id:274500 sequences / Arvadi Biofilm stones freshwater spring containing (among others) hydr AB425
FJ484024.1 gg_id:537733 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4840
GU208262.1 gg_id:553029 Study on Sediment Dongping Lake sediment clone 6F2-12 GU208
AY510250.1 gg_id:109312 microbial mat sulfidic cave spring Lower Kane Cave clone LKC3_198.8 AY510
AB176697.1 gg_id:107729 subsurface geothermal water clone HAuD-LB/2-3 AB176
FJ901811.1 gg_id:538289 phreatic limestone sinkholes northeastern Mexico water column surface cenote La Pa FJ9018
EU037198.1 gg_id:249812 Hot Spring Microbial Mat Khir Ganga India hot spring microbial mat clone MO43 EU037
DQ837278.1 gg_id:255039 Archaeal and pristine National Park Spain Donana coastal aquifer clone 56S_1B_81 DQ837
AB176695.1 gg_id:107438 subsurface geothermal water clone HAuD-MB/2-1 AB176
FJ901837.1 gg_id:558734 phreatic limestone sinkholes northeastern Mexico water column surface cenote La Pa FJ9018
AY955086.1 gg_id:114743 sediment clone 5-25R AY955
DQ145970.1 gg_id:138498 Thiobacillus sp. ML2-16 DQ145
DQ145974.1 gg_id:138290 Thiobacillus sp. ML2-40 DQ145
CU924728.1 gg_id:563027 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU924
AY510243.1 gg_id:109305 microbial mat sulfidic cave spring Lower Kane Cave clone LKC3_102B.25 AY510
DQ145968.1 gg_id:151278 Thiobacillus sp. ML1-72 DQ145
U58019.1 gg_id:6772 Thiobacillus aquaesulis U5801
AY845252.1 gg_id:110030 Spirillum kriegii str. D-430 AY845
FJ901825.1 gg_id:512862 phreatic limestone sinkholes northeastern Mexico water column surface cenote La Pa FJ9018
FJ484428.1 gg_id:553673 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4844
DQ145964.1 gg_id:138129 Thiobacillus sp. ML1-16 DQ145
FJ517027.1 gg_id:569792 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled water c FJ5170
AJ536815.1 gg_id:110737 magnesite mine clone cMM319-05 AJ5368
DQ837267.1 gg_id:248472 Archaeal and pristine National Park Spain Donana coastal aquifer clone 56S_1B_30 DQ837
EF492934.1 gg_id:247725 Iron-manganese Nodule and Its Surrounding Soil soil around iron-manganese nodule EF4929
EF173352.1 gg_id:186982 Options situ remediation contaminated mixture chlorinated hydrocarbons hydrocarb EF1733
AJ316618.1 gg_id:64665 Thiobacillus plumbophilus str. DSM 6690 Gro7 AJ3166
DQ984604.1 gg_id:242053 LY oil-contaminated soil clone LYC068 DQ984
AM490714.1 gg_id:539260 Linking and functional nutrient spiraling mats (USA) microbial mat sulfidic cave sprir AM490
GQ090471.1 gg_id:416305 Topographical and Temporal Human Skin Microbiome skin popliteal fossa clone nbw GQ090
DQ837266.1 gg_id:238477 Archaeal and pristine National Park Spain Donana coastal aquifer clone 56S_1B_20 DQ837
GQ388880.1 gg_id:535635 quality deterioration drinking system during serious red water outbreak clone Z4 GQ388
FJ437813.1 gg_id:559929 Benthic okenone production Fayetteville New York Green Lake chemocline 20.5 m wa FJ4378
DQ837229.1 gg_id:243280 Archaeal and pristine National Park Spain Donana coastal aquifer clone 49S1_1B_49 DQ837
EU735705.1 gg_id:355524 Biological Arsenic Remediation Project Groundwater Aquifer soil clone 1As11 EU735
DQ415761.1 gg_id:157821 Dominant microbial limestone-corroding biofilms frasassi system Italy Frasassi sulfid DQ415
GU208472.1 gg_id:581588 Study on Sediment Dongping Lake sediment clone Shc5-21 GU208
DQ520198.1 gg_id:160396 shal China sequences hypertrophic freshwater lake clone ML-9-101.2 DQ520

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DQ066957.1 gg_id:130688 Lake Washington sediment clone pLW-10 DQ066
GQ472317.1 gg_id:550530 Microbial Dongping Lake lake sediment clone 2B3-20 GQ472
EU360497.1 gg_id:269449 pilot biological ferrous iron oxidation mine water treatment plant clone W-Z-54 EU3604
FJ484433.1 gg_id:551992 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4844
AY532537.1 gg_id:104052 uranium-contaminated aquifer clone 1013-1-CG8 AY5325
AF407390.1 gg_id:19644 MCB-contaminated groundwater-treating reactor clone Ra9C4 AF4073
FJ712609.1 gg_id:516574 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) K FJ7126
GU208482.1 gg_id:589837 Study on Sediment Dongping Lake sediment clone Shc1-3 GU208
AB478665.1 gg_id:543991 SSU sequences Alps Cadagno di Fuori submerged biofilm mats montane wetland clo AB478
AY212734.1 gg_id:99462 water 20 m upstream manure clone 3up AY2127
EF133508.1 gg_id:175927 Ferribacter polymyxa str. PSTR EF1335
AJ519624.1 gg_id:74988 Uranium mill tailings soil sample clone Sh765B-TzT-132 proteobacterium AJ5196
EU340171.1 gg_id:245116 on Aquatic Macrophytes Geothermally Active Area Sedge Bay Yellowstone Lake WY EU3401
EU676421.1 gg_id:297924 rhizosphere pioneer plants (Bahia xylopoda and Viguiera linearis) growing on metals EU6764
EU676422.1 gg_id:290525 rhizosphere pioneer plants (Bahia xylopoda and Viguiera linearis) growing on metals EU6764
AJ536818.1 gg_id:109678 magnesite mine clone cMM319-36 AJ5368
AF445741.1 gg_id:41556 travertine hot spring clone SM2G09 AF4457
NR_025556.1 gg_id:569678 Tepidiphilus margaritifer str. N2-214; DSM 15129 NR_025
DQ287787.1 gg_id:143113 Methylobacillus flagellatus str. KT DQ287
AY298905.2 gg_id:88207 Methylobacillus pratensis str. F31 AY2989
AY027801.1 gg_id:42693 Aminomonas aminovorus str. C2A1; NCIMB 11268 AY0278
EU194898.1 gg_id:358307 Methylobacillus sp. str. M8 EU1948
AY772089.1 gg_id:132690 Methylophilus quaylei str. MT AY7720
AB193725.1 gg_id:108436 Methylophilus leisingeri str. DSM 6813 AB1937
AB193724.1 gg_id:108435 Methylophilus methylotrophus str. NCIMB 10515 AB1937
DQ287786.1 gg_id:141866 Methylotenera mobila str. JLW8 DQ287
FM872938.1 gg_id:366699 occupant source house floor dust clone FB04H04 FM872
FJ626263.1 gg_id:545204 Microbial larval insect gut field and laboratory sugarcane weevil Sphenophorus (Colec FJ6262
EU340223.1 gg_id:251118 on Aquatic Macrophytes Geothermally Active Area Sedge Bay Yellowstone Lake WY EU3402
AY436794.1 gg_id:104555 Methylophilus sp. str. ECd5 AY4367
AY546500.1 gg_id:99740 soda lake sediment clone 1H_31 AY5465
EF192877.1 gg_id:202563 sediments temperate artificial lake Rapel reservoir sediment clone Rap1_6A EF1928
DQ450168.1 gg_id:153929 (Uruguay) Atlantic coastal lagoon clone R7C30 DQ450
AY948052.1 gg_id:143623 Synchrony and seasonality two temperate rivers USA: Massachusetts Parker River cl AY9480
DQ228415.1 gg_id:142498 microbial compositions two subglacial possible role microbes chemical weathering p DQ228
FJ444649.1 gg_id:511215 Cotton Rhizosphere cotton rhizosphere clone 2h-19 FJ4446
EU662580.1 gg_id:533328 Productivity- Chemolithoautotrophically Based Sulfidic Karst Systems floating microb EU6625
AY947940.1 gg_id:143558 Synchrony and seasonality two temperate rivers USA: Massachusetts Ipswich River c AY9479
EU537789.1 gg_id:285536 profile human microbiota antecubital fossa (inner elbow) skin clone nbt79g11 EU5377
EU340172.1 gg_id:250544 on Aquatic Macrophytes Geothermally Active Area Sedge Bay Yellowstone Lake WY EU3401
EU801545.1 gg_id:306546 It's all ranking aquatic Chesapeake Bay MD clone 3C002863 EU8015

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AY212692.1 gg_id:105727 water 10 m downstream manure clone 244ds10	AY2126
EF580978.1 gg_id:278346 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM1	EF5809
EF019146.1 gg_id:206079 aspen rhizosphere clone Amb_16S_1787	EF0191
EU703371.1 gg_id:396964 lakes on Eastern Tin Plateau freshwater lake altitude 4298 masl clone XZTSH19	EU7033
AB464941.2 gg_id:343674 Altitudinal changes flora on Gulkana Glacier Alaska analyzed sequencing and -cycle c	AB4649
AY486132.1 gg_id:103599 Methylovorus mays str. C	AY4861
EU268114.1 gg_id:250802 summer Dalian and Liaodong bay determined sea water clone ZL1702	EU2681
EU799457.1 gg_id:330088 It's all ranking aquatic Newport Harbour RI clone 1C227059	EU7994
EU287055.1 gg_id:235281 Pacific arctic surface sediment clone B78-91	EU2870
EU591642.1 gg_id:264560 Microbial Ecology Focused-Pulsed Electric Field Sludge Pretreatment Methanogenic	EU5916
EU801464.1 gg_id:308817 It's all ranking aquatic Chesapeake Bay MD clone 3C002760	EU8014
AY238979.1 gg_id:105790 sludge dentrifying reactor fed with methanol clone DSBR-C043	AY2389
EU804023.1 gg_id:338998 It's all ranking aquatic Lake Gatun clone 5C231706	EU8040
EU236355.1 gg_id:289508 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg91B9	EU2363
DQ513082.1 gg_id:161213 Microbial life fluids ridge flank crustal fluid clone FS266-46B-03	DQ513
GQ350653.1 gg_id:543211 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich	GQ350
DQ066956.1 gg_id:135981 Lake Washington sediment clone pLW-13	DQ066
AJ400352.1 gg_id:29664 marine clone ZD0412	AJ4003
EF492937.1 gg_id:240233 Iron-manganese Nodule and Its Surrounding Soil soil around iron-manganese nodule	EF4929
AY678517.1 gg_id:103273 estuarine sediment clone C319a-R8C-D8	AY6785
EU801551.1 gg_id:329876 It's all ranking aquatic Chesapeake Bay MD clone 3C002872	EU8015
AY360530.1 gg_id:137280 Stable isotope probing dynamic methylotroph trophic interactions fungi and protozo	AY3605
FJ612340.1 gg_id:548303 and Biogeochemistry Dongping Lake Ecosystems lake water clone DP10.2.8	FJ6123
AY360525.1 gg_id:137243 Stable isotope probing dynamic methylotroph trophic interactions fungi and protozo	AY3605
GQ340339.1 gg_id:568552 Prokaryotic Marathonas Reservoir Greece water column clone VE08-118-BAC	GQ340
DQ810320.1 gg_id:170033 Oxygen Minimum Zone Surface- Lower Waters and Other Marine oxygen minimum	DQ810
AY546509.1 gg_id:101169 soda lake sediment clone 5H_52	AY5465
EU804036.1 gg_id:306270 It's all ranking aquatic Lake Gatun clone 5C231720	EU8040
AB294945.1 gg_id:219027 -dependent & analyses hydrothermal system off Taketomi Island Japan microbial ma	AB2949
EU800826.1 gg_id:328759 It's all ranking aquatic Delaware Bay NJ clone 2C229045	EU8008
EU037912.1 gg_id:247600 Metabolic Profiles Microbial Chlorinated Pesticides Contaminated Sites Three Geogr	EU0379
AY360533.1 gg_id:137226 Stable isotope probing dynamic methylotroph trophic interactions fungi and protozo	AY3605
EU800277.1 gg_id:338875 It's all ranking aquatic Delaware Bay NJ clone 2C228317	EU8002
GQ350509.1 gg_id:580430 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich	GQ350
GQ472336.1 gg_id:560126 Microbial Dongping Lake lake sediment clone 2S4-17	GQ472
GU230465.1 gg_id:587327 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A	GU230
AY792265.1 gg_id:125742 Humic Lake clone CrystalBog021G5	AY7922
EU287027.1 gg_id:255894 Pacific arctic surface sediment clone B78-63	EU2870
EF173353.1 gg_id:177340 Options situ remediation contaminated mixture chlorinated hydrocarbons hydrocarb	EF1733
FJ612406.1 gg_id:584989 and Biogeochemistry Dongping Lake Ecosystems lake water clone DP10.4.19	FJ6124

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EU803303.1 gg_id:332100 It's all ranking aquatic Lake Gatun clone 5C230853 EU803
FJ826390.1 gg_id:567795 Succession during spring diatom bloom Sea filtered surface sea water contrasting nor FJ8263
AJ318162.1 gg_id:24165 Waste-gas biofilter clone BIqi38 AJ3181
EF379433.1 gg_id:212747 sea water Victoria Harbor (Hong Kong) area clone TLC-PA3-8 EF3794
GU183586.1 gg_id:580735 Structures Reduced and Oxidized Aquifer lower Aqueous Arsenic Titas Bangladesh a GU183
AJ582036.1 gg_id:93630 soil near uranium mill tailings clone JG36-GS-101 AJ5820
FJ612167.1 gg_id:587500 and Biogeochemistry Dongping Lake Ecosystems lake water clone DP3.4.51 FJ6121
EU803922.1 gg_id:303542 It's all ranking aquatic Lake Gatun clone 5C231580 EU8039
EF516180.1 gg_id:222196 grassland soil clone FCPU667 EF5161
AY360552.1 gg_id:137233 Stable isotope probing dynamic methylotroph trophic interactions fungi and protozo AY3605
AF328143.1 gg_id:38266 Simonsiella crassa str. ATCC29447 AF3281
AF487710.1 gg_id:55486 Alysiella filiformis str. ATCC 15532 AF4877
EU661705.1 gg_id:282842 Aquitalea magnusonii str. NBRAJG85 EU6617
EU661703.1 gg_id:284709 Aquitalea magnusonii str. NBRAJG 83 EU6617
EU661704.1 gg_id:285068 Aquitalea magnusonii str. NBRAJG84 EU6617
EU741686.1 gg_id:332151 Aquitalea sp. str. LT10-3 EU7416
GQ916513.1 gg_id:554457 Aquitalea magnusonii str. AOLR15 GQ916
EU548073.1 gg_id:276722 Aquitalea magnusonii str. 5YN1-11 EU5480
AB087265.2 gg_id:63985 Neisseria denitrificans str. IAM 14975 AB0872
AY117557.1 gg_id:63496 Chromobacterium violaceum str. 07 AY1175
AE016825.1 gg_id:148906 Chromobacterium violaceum str. ATCC 12472 AE0168
AY117559.1 gg_id:63494 Chromobacterium sp. str. 46 AY1175
AY117571.1 gg_id:63483 Chromobacterium sp. str. 70 AY1175
EU372837.1 gg_id:270823 Chromobacterium violaceum str. ESBV 4400 EU3728
AY117552.1 gg_id:63505 Chromobacterium sp. str. 48 AY1175
EU693450.1 gg_id:298487 Chromobacterium violaceum str. EAV2 EU6934
AB426118.1 gg_id:262569 Chromobacterium sp. str. DS-1 AB4261
AY117565.1 gg_id:63489 Chromobacterium sp. str. 53 AY1175
AY344056.1 gg_id:101677 Chromobacterium suttsuga str. PRAA4-1 AY3440
AF328152.1 gg_id:33892 Simonsiella sp. 27381 str. ATCC 27381; ICPB 3601 ATCC AF3281
GQ114060.1 gg_id:378482 Topographical and Temporal Human Skin Microbiome skin interdigital web space clc GQ114
AF328151.1 gg_id:21796 Simonsiella sp. 29465 str. ATCC 29465; ICPB 3648 ATCC AF3281
GQ114732.1 gg_id:427571 Topographical and Temporal Human Skin Microbiome skin interdigital web space clc GQ114
AF328154.1 gg_id:73115 Simonsiella steedae str. ATCC29435 AF3281
AB087261.2 gg_id:56599 Simonsiella steedae str. IAM 14972 AB0872
AF328155.1 gg_id:21461 Simonsiella steedae str. ATCC29445 AF3281
AM397080.1 gg_id:170117 Deefgea rivuli str. WB 3.4-79 AM397
FJ347759.1 gg_id:359570 Deefgea sp. str. nsw-4 FJ3477
AB525415.1 gg_id:575021 Eikenella corrodens str. JCM 12952 AB5254
GQ422740.1 gg_id:574200 Eikenella corrodens str. Moore D25 GQ422
EF509092.1 gg_id:226450 Loss During Antibiotic Treatment 1 Intubated Patients Colonized Pseudomonas aerug EF5090

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NZ_ACJW02000005.1 gg_id:470523 Kingella oralis str. ATCC 51147 NZ_AC
AY644511.1 gg_id:107215 Kingella sp. M03-232 AY6445
AY349388.1 gg_id:92601 human oral cavity clone JC012 AY3493
DQ003616.1 gg_id:125235 Kingella genomosp. P1 oral cone MB2_C20 DQ003
GU227231.1 gg_id:536311 Microbiological immunological Adolescent Moroccan Aggressive Periodontitis Patier GU227
EU118276.2 gg_id:248652 Laribacter hongkongensis str. LHHZ69f EU118
U89333.1 gg_id:6367 Microvirgula aerodenitrificans str. Sgly2 U89333
DQ117531.1 gg_id:133808 Neisseria bacilliformis str. MDA1552 DQ117
AY560519.1 gg_id:98975 Neisseria bacilliformis str. MDA2833 AY5605
DQ117530.1 gg_id:134552 Neisseria bacilliformis str. CCUG38158 DQ117
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AJ239302.1 gg_id:6408 Neisseria elongata str. I3 AJ2393
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AF398320.1 gg_id:62759 Neisseria meningitidis str. M8237 AF3983
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DQ201330.1 gg_id:140124 Neisseria meningitidis str. CMCC(B)29034 DQ201
AF382302.1 gg_id:65958 Neisseria meningitidis str. M7855 AF3823
AY146978.1 gg_id:77115 Neisseria meningitidis str. N.451/91 AY1469
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AF310541.1 gg_id:32179 Neisseria meningitidis str. M239 AF3105
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AF409943.1 gg_id:60511 Neisseria meningitidis str. M1141 AF4099
AF310406.1 gg_id:73554 Neisseria meningitidis str. M7153 AF3104
AF398321.1 gg_id:62758 Neisseria meningitidis str. M8236 AF3983
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AF409944.1 gg_id:60507 Neisseria meningitidis str. M2671 AF4099
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AJ239312.1 gg_id:6471 Neisseria meningitidis str. S3446 AJ2393
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AF479578.1 gg_id:16823 Neisseria subflava str. NJ9703 AF4795
L10738.1 gg_id:6396 Neisseria weaveri L10738
AF395911.1 gg_id:51956 Neisseria weaveri VA6362grgr_2001 AF3959
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AF310565.1 gg_id:32023 Neisseria meningitidis str. M2786 AF3105
GU233442.1 gg_id:517381 Neisseria iguanae str. NVSL 85737 GU233
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FJ191574.1 gg_id:335421 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1915
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AF398308.1 gg_id:62770 Neisseria meningitidis str. M7509 AF3983

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DQ201336.1 gg_id:140609 Neisseria meningitidis str. CMCC(B)29202 DQ201
FJ191449.1 gg_id:316460 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1914
AF382307.1 gg_id:65953 Neisseria meningitidis str. M7925 AF3823
AF310344.1 gg_id:30575 Neisseria meningitidis str. M3291 AF3103
EF509732.1 gg_id:218613 Loss During Antibiotic Treatment 1 Intubated Patients Colonized Pseudomonas aerug EF5097
AF310551.1 gg_id:52834 Neisseria meningitidis str. M4797 AF3105
AF398275.1 gg_id:62802 Neisseria meningitidis str. M812 AF3982
AF409938.1 gg_id:60518 Neisseria meningitidis str. M4644 AF4099
FJ191958.1 gg_id:321506 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1919
DQ201323.1 gg_id:140409 Neisseria meningitidis str. CMCC(B)29019 DQ201
DQ201329.1 gg_id:140166 Neisseria meningitidis str. CMCC(B)29031 DQ201
AJ239284.1 gg_id:6465 Neisseria lactamica str. L20 AJ2392
AY132119.1 gg_id:59602 Neisseria meningitidis str. M8574 AY1321
FJ191979.1 gg_id:350810 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1919
AJ239293.1 gg_id:6442 Neisseria sicca str. Q28 AJ2392
AJ239280.1 gg_id:6428 Neisseria flavescens str. LNP444 AJ2392
FJ192352.1 gg_id:350507 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1923
FJ976263.1 gg_id:539814 Interindividual healthy human oral microbiota mouth clone 6BBB59 FJ9762
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NZ_ACQV01000025.1 gg_id:470744 Neisseria flavescens str. SK114 NZ_AC
FJ192317.1 gg_id:318935 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1923
AF398273.1 gg_id:62803 Neisseria meningitidis str. M1168 AF3982
AF398301.1 gg_id:62777 Neisseria meningitidis str. M7918 AF3983
FJ192355.1 gg_id:353666 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1923
DQ201334.1 gg_id:139856 Neisseria meningitidis str. CMCC(B)29046 DQ201
AY648034.1 gg_id:99627 Neisseria meningitidis str. NZ00/215 AY648
AF398325.1 gg_id:62755 Neisseria meningitidis str. N.33/00 AF3983
NZ_ACDX01000110.1 gg_id:470729 Neisseria mucosa str. ATCC 25996 NZ_AC
L06171.1 gg_id:6402 Neisseria elongata L06171
AF409941.1 gg_id:72478 Neisseria meningitidis str. M1417 AF4099
AF398252.1 gg_id:62824 Neisseria meningitidis str. M5283 AF3982
AB194302.1 gg_id:108299 Sivamonas terrae str. KM-45 AB1943
AF328146.1 gg_id:69998 Simonsiella muelleri str. ATCC29441 AF3281
FJ976380.1 gg_id:554114 Interindividual healthy human oral microbiota mouth clone 22B524 FJ9763
GQ102572.1 gg_id:447914 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ102
DQ003630.1 gg_id:126626 human mouth clone MB5_P15 DQ003
GQ106230.1 gg_id:468425 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ106
EF212919.1 gg_id:214313 Stenoxybacter acetivorans str. TAM-WH4 EF2129
DQ837616.1 gg_id:164644 structures honeybee intestines and their response insecticidal proteins gut clone HB DQ837
AB192065.1 gg_id:144630 termite gut homogenate clone M1NP1-30 AB1920

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EU370420.1 gg_id:264994 Neisseria sp. str. 93S1 EU370
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GQ068667.1 gg_id:466523 Topographical and Temporal Human Skin Microbiome skin external auditory canal cl GQ068
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FJ502346.1 gg_id:512007 Neisseria sp. str. GRW38 FJ5023
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GQ081072.1 gg_id:384019 Topographical and Temporal Human Skin Microbiome skin external auditory canal cl GQ081
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GQ068787.1 gg_id:376358 Topographical and Temporal Human Skin Microbiome skin external auditory canal cl GQ068
AJ239288.1 gg_id:6399 Neisseria animalis str. NCTC10212 AJ2392
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Y17602.1 gg_id:6368 Formivibrio citricus str. DSM 6150 Y17602

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FJ470512.1 gg_id:540352 Chinese human cavities healthy oral cavity clone A_D_03_77 FJ4705
GQ131417.1 gg_id:390069 Neisseria sp. oral taxon 014 str. F0314 GQ131
GQ081262.1 gg_id:401550 Topographical and Temporal Human Skin Microbiome skin external auditory canal cl GQ081
GQ093550.1 gg_id:374530 Topographical and Temporal Human Skin Microbiome skin glabella clone nbw366b1 GQ093
GQ068928.1 gg_id:426161 Topographical and Temporal Human Skin Microbiome skin external auditory canal cl GQ068
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AF487709.1 gg_id:58031 Neisseria dentiae str. V33 AF487
GQ041790.1 gg_id:437690 Topographical and Temporal Human Skin Microbiome skin glabella clone nbw932f05 GQ041
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GQ081230.1 gg_id:462985 Topographical and Temporal Human Skin Microbiome skin external auditory canal cl GQ081
FJ593907.1 gg_id:574480 Chitinibacter sp. str. TNR-14 FJ5939
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DQ376554.1 gg_id:151309 Microbial ecology removing phosphate wastewater EBPR process aerobic sequencin DQ376
GQ080988.1 gg_id:456420 Topographical and Temporal Human Skin Microbiome skin external auditory canal cl GQ080
AJ629192.1 gg_id:129475 Kingella potus str. 3/SID/1128 AJ6291
DQ006842.1 gg_id:118296 Neisseria sp. R-24680 DQ006
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GQ081261.1 gg_id:446559 Topographical and Temporal Human Skin Microbiome skin external auditory canal cl GQ081
AY005028.1 gg_id:53569 Neisseria sp. oral strain str. B33KA AY005
DQ314581.1 gg_id:148882 chitin-degrading Taiwan C14 DQ314
AY609199.2 gg_id:140478 Chromobacterium sp. str. 2002 AY609
EU536080.1 gg_id:280825 profile human microbiota antecubital fossa (inner elbow) skin clone nbt08d11 EU536
AB013258.1 gg_id:6372 deep-sea sediment clone NKB6 AB013
AM295491.1 gg_id:164660 Gulbenkianus mobilis str. E4FC31 AM295
AY904024.1 gg_id:146009 Aquaspirillum sp. str. 411 AY904

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AB286476.1 gg_id:206343 Reducing production and domination comamonadaceae reducing oxygen supply was AB286
AB076000.1 gg_id:64884 Aquaspirillum putridiconchylium str. IAM 14964 AB076
FJ375419.1 gg_id:589312 T-RFLP powerful tool rapid assessment cells microbial fuel cell sheet carbon cloth ano FJ3754
L06174.1 gg_id:6375 Vitreoscilla stercoraria L06174
AF227863.1 gg_id:36932 str. 82348 AF227
AM689950.1 gg_id:226696 Vogesella sp. str. SK-2 AM689
AF287297.1 gg_id:29792 Nitrosococcus mobilis str. Nc2 AF287
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AF272421.1 gg_id:38503 Nitrosomonas sp. NM 41 AF272
AJ621028.1 gg_id:105866 Nitrosomonas sp. Is86 AJ6210
AF272413.1 gg_id:40104 Nitrosomonas halophila AF272
FJ529930.1 gg_id:542891 Initial ammonia- and nitrite-oxidizing compromises successful nitritation counter-diffu FJ5299
AJ621032.1 gg_id:101077 Nitrosomonas sp. Is343 AJ6210
AF363289.1 gg_id:40156 Nitrosomonas sp. str. K1 AF363
EF092247.1 gg_id:265873 Persistent Axinella corrugata Nitrosomonas clone B276_F5 EF0922
AF272418.1 gg_id:16711 Nitrosomonas marina AF272
AF353164.1 gg_id:22792 Nitrosomonas sp. DYS323 AF353
AY123794.1 gg_id:69852 Nitrosomonas sp. str. Nm143 AY123
GQ396862.1 gg_id:559849 Functional unvegetated perhumid recently-deglaciated soils do not correlate shifts s GQ396
EF016119.1 gg_id:182102 Nitrosomonas oligotropha str. AS1 EF0161
EF092216.1 gg_id:268524 Persistent Axinella corrugata Nitrosomonas clone B2505_H7 EF0922
GU183580.1 gg_id:583485 Structures Reduced and Oxidized Aquifer lower Aqueous Arsenic Titas Bangladesh a GU183
AY123795.1 gg_id:69328 Nitrosomonas eutropha str. Nm57 AY123
AY123807.1 gg_id:70906 Nitrosospira multiformis str. Nl13; C-71; ATCC 25196 AY123
AY856377.1 gg_id:110754 Nitrosospira sp. str. TCH711 AY856
AY856375.1 gg_id:109672 Nitrosospira sp. str. HBN8222A AY856
AF353156.1 gg_id:20624 Nitrosospira sp. TCH716 AF353
AY123804.1 gg_id:72266 Nitrosospira sp. str. Nsp17 AY123
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X84662.1 gg_id:6504 Nitrosospira sp. X84662
AY123803.1 gg_id:69463 Nitrosovibrio tenuis str. Nv1 AY123
M96405.1 gg_id:6602 Nitrosovibrio tenuis str. Nv12 M9640
AY123800.1 gg_id:70700 Nitrosospira briensis str. Nsp10 AY123
AY631269.1 gg_id:102482 Nitrosovibrio sp. FJI82 AY631
EU050920.1 gg_id:250427 libraries sediment Kings Bay Svalbard Arctic clone SS1_B_07_05 EU050
AF414581.1 gg_id:33825 uranium mine sediment clone P3OB-51 AF414
AF080256.1 gg_id:6535 Nitrosospira sp. str. TYM9 AF080
AJ630749.1 gg_id:105991 marine surface water clone BET_1D11G67 AJ6307
EU734961.1 gg_id:311415 assemblages sediment station DBSE Northern Bering Sea clone 054D10 EU734
AY123813.1 gg_id:73969 Nitrosospira sp. str. Nsp65 AY123
L35505.1 gg_id:6517 Nitrosospira briensis str. C-128 L35505

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AF363291.1 gg_id:40900 Nitrosovibrio sp. str. RY6A AF3632
EU287131.1 gg_id:245816 Pacific arctic surface sediment clone P13-38 EU2871
X90820.1 gg_id:6538 Nitrosospira sp str. AHB1 X90820
AJ005545.1 gg_id:6518 Nitrosospira sp. F3 AJ0055
AY631270.1 gg_id:106086 Nitrosovibrio sp. FJI423 AY6312
AF363290.1 gg_id:43839 Nitrosovibrio sp. str. RY3C AF3632
M96396.1 gg_id:6503 Nitrosospira briensis M9639
GQ396891.1 gg_id:592669 Functional unvegetated perhumid recently-deglaciated soils do not correlate shifts s GQ396
EU676432.1 gg_id:288399 rhizosphere pioneer plants (Bahia xylopoda and Viguiera linearis) growing on metals- EU6764
AF177427.1 gg_id:37585 Acanthamoeba sp. UWC12 symbiont AF1774
AF177425.1 gg_id:43515 Acanthamoeba sp. Page 23 symbiont AF1774
AF352385.1 gg_id:29264 Acanthamoeba sp. TUMSJ-226 symbiont AF3523
DQ280369.1 gg_id:144094 Microvirgula sp. str. HW7 DQ280
EU708500.1 gg_id:296975 and Situ Quantification Arsenite Oxidizing Denitrifying Enrichment anoxic arsenite ox EU7085
AF331975.1 gg_id:66610 Azoarcus sp. str. HxN1 AF3319
Y13222.1 gg_id:6771 Azoarcus sp. str. 22Lin Y13222
NR_024970.1 gg_id:543017 Azoarcus toluclasticus str. MF63; ATCC 700605 NR_024
AF123075.1 gg_id:6778 Azoarcus tolulyticus str. BL11 AF1230
AF123076.1 gg_id:6779 Azoarcus toluclasticus str. MF7 AF1230
L33687.1 gg_id:6804 Azoarcus tolulyticus str. Td-1 L33687
L33693.1 gg_id:6782 Azoarcus tolulyticus str. Td-3 L33693
L33692.1 gg_id:6784 Azoarcus tolulyticus str. Td-21 L33692
DQ264504.1 gg_id:141528 universal microarray broader than typical library when sampling subsurface groundv DQ264
DQ264480.1 gg_id:142574 universal microarray broader than typical library when sampling subsurface groundv DQ264
L33688.1 gg_id:6766 Azoarcus tolulyticus str. Td-15 L33688
AB234249.1 gg_id:148299 Biotransformation Polychlorinated Dioxins Microbial Dynamics Sediment Microcosm AB2342
DQ264419.1 gg_id:143835 universal microarray broader than typical library when sampling subsurface groundv DQ264
L33689.1 gg_id:6781 Azoarcus tolulyticus str. Td-17 L33689
DQ264565.1 gg_id:142083 universal microarray broader than typical library when sampling subsurface groundv DQ264
AF229861.1 gg_id:35148 Azoarcus tolulyticus str. 2FB6 AF2298
EU528179.1 gg_id:286700 Seasonal Dynamics Mudflat Mouth Major Kentucky Lake Reservoir Tributary sedime EU5281
AJ630292.1 gg_id:100647 Azonexus fungiphilus AJ6302
GU056304.1 gg_id:557513 Anoxic Oxidation Arsenite Arsenate linked Chlorate Reduction returned actived slud GU056
AF011347.1 gg_id:6714 Azospira oryzae str. 6a3 AF0113
AY235689.1 gg_id:106890 Nitrogen-fixation strains wild rice AY2356
AY171616.1 gg_id:77627 Dechlorosoma sp. str. AH AY1716
DQ863512.1 gg_id:165690 Azospira oryzae str. N1 DQ863
AF170348.1 gg_id:6722 Azospira oryzae str. PS AF1703
AY277622.1 gg_id:101136 Azospira oryzae str. GR-1 AY2776
FJ391490.1 gg_id:367951 primers targeting Gallionella- neutrophilic oxidizers enrichments and soils iron rich irr FJ3914
AF011349.1 gg_id:6675 Azovibrio sp. str. BS20-3 AF0113

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FJ535057.1 gg_id:591896 Enhancement protein conversion volatile fatty acids accumulation during addition: ef FJ5350
DQ640664.1 gg_id:233934 full-scale EBPR sludge clone Skagenf42 DQ640
GQ340232.1 gg_id:551430 Prokaryotic Marathonas Reservoir Greece water column clone VE07-68-BAC GQ340
AM084133.1 gg_id:140097 Dechloromonas sp. str. R-28400 AM084
AB240507.1 gg_id:142587 microbial structure rhizosphere biofilm Sapporo root-base (80 160 mm) Phragmites S AB240
FJ719082.1 gg_id:531877 Protozoan grazers alter metabolically active aquifer aquifier sediments clone p04_E0 FJ7190
EF515635.1 gg_id:224951 Electricigen Enrichment MFC full-scale anaerobic bioreactor sludge treating brewery EF5156
DQ376553.1 gg_id:153255 Microbial ecology removing phosphate wastewater EBPR process aerobic sequencin DQ376
CU466914.1 gg_id:319502 Evry wastewater treatment plant anoxic basin clone CU466
CU920167.1 gg_id:572334 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU920
CU466920.1 gg_id:304463 Evry wastewater treatment plant anoxic basin clone CU466
AB240296.1 gg_id:144181 microbial structure Phragmites rhizosphere biofilm reed bed reactor laboratory clon AB240
CU466858.1 gg_id:344521 Evry wastewater treatment plant anoxic basin clone CU466
FJ444762.1 gg_id:542590 Cotton Rhizosphere cotton rhizosphere clone 4y-107 FJ4447
CU926684.1 gg_id:554053 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU926
Y17060.1 gg_id:6706 Ferribacterium limneticum str. cda-1 Y17060
DQ296471.1 gg_id:161279 Microbial dissection quinoline-degradation lab-scale anaerobic biofilm reactor quino DQ296
CU466895.1 gg_id:319055 Evry wastewater treatment plant anoxic basin clone CU466
DQ923115.1 gg_id:163080 Methyloversatilis universalis str. 500 DQ923
FJ167509.1 gg_id:333442 microbiology granular sludge under autotrophic mixotrophic and heterotrophic denit FJ1675
FJ902588.1 gg_id:568555 phreatic limestone sinkholes northeastern Mexico biofilm 1m deep hydrothermal ver FJ9025
FM207922.1 gg_id:305052 Anaerobic p-xylene mineralization activated sludge and fresh water sediment denitr FM207
DQ088734.1 gg_id:137170 subsurface clone BE16FW031601GDW_hole1-11 DQ088
AB234251.1 gg_id:150027 Biotransformation Polychlorinated Dioxins Microbial Dynamics Sediment Microcosm AB234
DQ676301.1 gg_id:224877 Archaeal sediment and plankton freshwater pond suboxic freshwater-pond clone M DQ676
FM207901.1 gg_id:328865 Anaerobic p-xylene mineralization activated sludge and fresh water sediment denitr FM207
DQ327692.1 gg_id:166580 Biological filtration limits carbon availability affects downstream biofilm formation a DQ327
AY945886.1 gg_id:146463 Thauera and Azoarcus functionally important genera denitrifying quinoline-removal AY945
FJ484883.1 gg_id:511775 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4848
DQ088748.1 gg_id:129402 subsurface clone BE325FW032701CTS_hole1-85 DQ088
FM207903.1 gg_id:327175 Anaerobic p-xylene mineralization activated sludge and fresh water sediment denitr FM207
FM207916.1 gg_id:351540 Anaerobic p-xylene mineralization activated sludge and fresh water sediment denitr FM207
EF363051.1 gg_id:268998 Microbial life within saline fluid Continental Deep Drilling German continental deep g EF3630
AB252920.1 gg_id:171974 Identification Shibayama lagoon Ishikawa Japan iron-oxidation biofilm clone 010 AB2529
AY879297.1 gg_id:111577 Denitratisoma oestradiolicum str. AcBE2-1 AY8792
FM207918.1 gg_id:344874 Anaerobic p-xylene mineralization activated sludge and fresh water sediment denitr FM207
FJ485121.1 gg_id:533665 microbial autonomous rover world's deepest phreatic sinkhole water column El Zacat FJ4851
DQ676390.1 gg_id:222171 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676
GQ340363.1 gg_id:538380 Prokaryotic Marathonas Reservoir Greece water column clone VE08-161-BAC GQ340
EF205275.1 gg_id:200507 Arctic gold mine fracture water clone Lupin-1130m-1-MDA-ubp1 EF2052

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AY235684.1 gg_id:106107 Nitrogen-fixation strains wild rice AY2356
AJ306683.1 gg_id:27286 Sterolibacterium denitrificans str. Chol-1S AJ3066
FJ391504.1 gg_id:369022 primers targeting Gallionella- neutrophilic oxidizers enrichments and soils iron rich irr FJ3915
DQ836749.1 gg_id:163176 attached biomass denitrifying bioreactor clone DB-5 DQ836
AM934804.1 gg_id:556892 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM934
FM207927.1 gg_id:325823 Anaerobic p-xylene mineralization activated sludge and fresh water sediment denitr FM207
FM207953.1 gg_id:304406 Anaerobic p-xylene mineralization activated sludge and fresh water sediment denitr FM207
DQ532120.1 gg_id:161420 Ultramicrobacter hongkongensis str. 5a1 DQ532
DQ676425.1 gg_id:226206 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676
AM268369.1 gg_id:191997 Functional Redundancy Versatility Rhodocyclales Activated Sludge: Improvement ar AM268
AB186837.1 gg_id:107618 polychlorinated dioxin dechlorinating microcosm clone TSBW06 AB1868
AY532566.1 gg_id:101212 uranium-contaminated aquifer clone 1013-28-CG13 AY5325
AF234757.1 gg_id:51803 sludge clone S40 AF2347
AF204243.1 gg_id:31185 P+ sludge clone GC24 AF2042
AY762627.1 gg_id:108150 gold mine clone DSBACT86 AY7626
AB378589.1 gg_id:271761 dominate denitrification nitrate- succinate-ammended stable istope probing 13C fra AB3785
EF590007.2 gg_id:228477 changes structure sediments Songhua River contaminated nitrobenzene -DGGE cloni EF5900
AF072922.1 gg_id:6533 nitrifying/denitrifying industrial activated sludge clone S28 S28 AF0729
AY328574.1 gg_id:89345 drinking water system simulator clone HOClCi25 AY3285
GU454941.1 gg_id:530591 Understanding short-chain fatty acids accumulation enhanced alkaline digestion: kir GU454
EF492892.1 gg_id:236435 Iron-manganese Nodule and Its Surrounding Soil soil around iron-manganese nodule EF4928
EU037206.1 gg_id:239439 Hot Spring Microbial Mat Khir Ganga India hot spring microbial mat clone MO91 EU0372
DQ984607.1 gg_id:235493 LY oil-contaminated soil clone LYC108 DQ984
FM204950.1 gg_id:547160 Prokaryote shifts and reductive dehalogenase redundancy response electron donor FM204
EU835438.1 gg_id:352426 structure and quorum sensing reverse osmosis RO membrane biofilm clone 2B09 EU8354
DQ464135.1 gg_id:157026 acid mine drainage clone D3-25 DQ464
FM207915.1 gg_id:349221 Anaerobic p-xylene mineralization activated sludge and fresh water sediment denitr FM207
AB179684.1 gg_id:106337 groundwater 0.2 m-captured fraction clone KNA6-EB04 AB1796
DQ988263.1 gg_id:190625 Microbial lab-scale A1-A2-O fixed biofilm system coking wastewater treatment with DQ988
AB252918.1 gg_id:167769 Identification Shibayama lagoon Ishikawa Japan iron-oxidation biofilm clone 195 AB2529
EF565160.1 gg_id:246262 'Candidatus accumulibacter' structure enhanced biological phosphorus removal sludg EF5651
FJ356026.1 gg_id:366769 Targeted FACS enrichment groups lab scale systems anaerobic:aerobic lab-scale EBPR FJ3560
GU250852.1 gg_id:578062 Anaerobic biodegradation branched hydrocarbon biomarker analog pristane waste GU250
D16210.1 gg_id:6712 Rhodocyclus tenuis str. SW18 D16210
D16209.1 gg_id:6713 Rhodocyclus tenuis str. DSM110 D16209
AY297807.1 gg_id:88575 waterlogged archaeological wood clone pACH91 AY2978
AF227840.1 gg_id:26639 str. 51885 AF2278
DQ640677.1 gg_id:230074 full-scale EBPR sludge clone Skagenf18 DQ640
AY706424.1 gg_id:100543 TCE-contaminated groundwater clone G1a AY7064
Y17601.1 gg_id:6721 Propionivibrio dicarboxylicus str. DSM 5885 Y17601
AB062831.1 gg_id:54073 termite gut homogenate clone BCf6-08 AB0628

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AF502231.1 gg_id:59337 EBPR sludge clone HP1A26 AF5022
AB240294.1 gg_id:143468 microbial structure Phragmites rhizosphere biofilm reed bed reactor laboratory clon AB2402
AB089103.1 gg_id:78240 termite gut homogenate clone Rs-K54 AB0891
NR_024855.1 gg_id:545495 Propionivibrio pelophilus str. asp 66; DSM 12018 NR_024
AY064179.1 gg_id:65492 EBPR sludge clone UCTN173 AY0641
AF255631.1 gg_id:36331 EBPR sludge clone Ebpr16 AF2556
DQ087192.1 gg_id:134476 clone HE018 DQ087
AY691423.1 gg_id:105089 Rhodocyclus sp. str. HOD 5 AY6914
AF229868.1 gg_id:18623 Thauera aromatica str. 4FB2 AF2298
AF229867.1 gg_id:42293 Thauera aromatica str. 4FB1 AF2298
AF229882.1 gg_id:69798 Thauera aromatica str. 3CB3 AF2298
AF123264.1 gg_id:6759 Thauera chlorobenzoica str. 3CB-1 AF1232
AF229887.1 gg_id:36677 Thauera aromatica str. 3BB1 AF2298
AJ315678.1 gg_id:25264 Thauera phenylacetica str. B4P AJ3156
AJ315677.1 gg_id:40990 Thauera aminoaromatica str. S2 AJ3156
U95176.1 gg_id:6763 Thauera aromatica str. T1 U95176
AY548757.1 gg_id:111927 Thauera sp. str. Cin34 AY5487
EU434525.1 gg_id:273528 Thauera phenylacetica str. b208 EU4345
AY098637.2 gg_id:71611 industrial-phenol-degrading community clone MM1 sp. AY0986
DQ018816.1 gg_id:131190 anaerobic swine lagoon clone C-147 DQ018
AJ505850.1 gg_id:58572 Thauera sp. str. PIV-1 AJ5058
EU434574.1 gg_id:265716 Thauera aromatica str. b66 EU4345
FJ900965.1 gg_id:513787 Polymorphism microbial reservoirs water cut stage oil field clone G-3 FJ9009
AM982629.2 gg_id:279478 microbial and slurry pig saw dust spent bedding clone PISD-AJG10 AM982
AB210048.1 gg_id:122645 mini plant Run 5 Nippon Steel Research Engineering Center clone nsmpV06 AB2100
CU466853.1 gg_id:346949 Evry wastewater treatment plant anoxic basin clone CU4668
AJ853609.1 gg_id:111043 landfill leachate clone GZKB116 AJ8536
AY823976.1 gg_id:145956 Investigation Acetate-Fed Denitrifying Microbial Stable Isotope Probing Full-Cycle an AY8239
X68491.1 gg_id:6756 Thauera selenatis str. Ax X68491
EF205256.1 gg_id:197834 Thauera sp. 'CJAOPY3 (T-II)' str. CJAOPY3 (T-II) EF2052
FJ825447.1 gg_id:569266 Microbial during start-up mesophilic biogas plant clone 112_BE3_37 FJ8254
AY823967.1 gg_id:145778 Investigation Acetate-Fed Denitrifying Microbial Stable Isotope Probing Full-Cycle an AY8239
AY570693.1 gg_id:108565 Thauera sp. Al7 AY5706
Y17590.1 gg_id:6761 Thauera mechernichensis str. TL1 Y17590
EU434467.1 gg_id:271936 Thauera aminoaromatica str. a241 EU4344
AY945863.1 gg_id:149463 Thauera and Azoarcus functionally important genera denitrifying quinoline-removal AY9458
AY945915.1 gg_id:145574 Thauera and Azoarcus functionally important genera denitrifying quinoline-removal AY9459
Y17591.1 gg_id:6762 Thauera selenatis str. ATCC 55363T Y17591
AJ005816.1 gg_id:6747 Thauera linaloolentis str. 47Lol AJ0058
AM268367.1 gg_id:179901 Functional Redundancy Versatility Rhodocyclales Activated Sludge: Improvement ar AM268
DQ018794.1 gg_id:133087 anaerobic swine lagoon clone C-38 DQ018

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GQ107403.1 gg_id:424310 Topographical and Temporal Human Skin Microbiome skin popliteal fossa clone nbw GQ107
EU104249.1 gg_id:335484 Evidence Comamonadaceae determinants settling performance activated sludge clo EU1042
AM268359.1 gg_id:196002 Functional Redundancy Versatility Rhodocyclales Activated Sludge: Improvement ar AM268
GQ093262.1 gg_id:442735 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw42 GQ093
AB192185.1 gg_id:144844 termite gut homogenate clone RsTz-39 AB1921
AY265878.1 gg_id:82517 perchlorate-reducing PMC AY2658
GQ100922.1 gg_id:463566 Topographical and Temporal Human Skin Microbiome skin retroauricular crease clor GQ100
GQ086556.1 gg_id:424472 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw29 GQ086
AY124797.1 gg_id:69212 Dechloromonas sp. str. LT-1 AY1247
DQ376566.1 gg_id:152042 Microbial ecology removing phosphate wastewater EBPR process aerobic sequencin DQ376
AB369190.1 gg_id:254556 Microbiological Assessment Circulation Mud Fluids During First Operation Drilling De AB3691
AF072921.1 gg_id:6799 nitrifying/denitrifying industrial activated sludge clone S23 S23 AF0729
AF072918.1 gg_id:6786 nitrifying/denitrifying industrial activated sludge clone S3 S3 AF0729
DQ640700.1 gg_id:231550 full-scale EBPR sludge clone Skagenf76 DQ640
FJ823914.1 gg_id:526353 Controlling accumulation fermentation inhibitors biorefinery recycle water microbial FJ8239
GQ024542.1 gg_id:485128 Topographical and Temporal Human Skin Microbiome skin palm clone nbw123e11c1 GQ024
AY947989.1 gg_id:142103 Synchrony and seasonality two temperate rivers USA: Massachusetts Parker River cl AY9479
AF072919.1 gg_id:6787 nitrifying/denitrifying industrial activated sludge clone H25 H25 AF0729
GQ093869.1 gg_id:399333 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ093
EU015084.1 gg_id:236679 Investigation estrogen-degrading bioreactors membrane bioreactor suspension and EU0150
EF648064.1 gg_id:241562 dynamics during produced water treatment aerobic activated sludge clone HB61 EF6480
EU133837.1 gg_id:240971 tallgrass prarie soil clone FFCH15186 EU1338
AY160865.1 gg_id:96448 termite intestinal tract clone COB P5-5 AY1608
AB024934.1 gg_id:21532 Sphingomonas sp. A1 AB0249
DQ836761.1 gg_id:167461 attached biomass denitrifying bioreactor clone DB-86 DQ836
FJ478739.1 gg_id:511268 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4787
DQ974114.1 gg_id:245301 Azospira restricta str. SUA2 DQ974
FJ374246.1 gg_id:362367 Inter- and intraspecific humivorous scarab beetle larvae (Pachnoda spp.) larval hindgu FJ3742
AF204246.1 gg_id:50481 EBPR sludge clone SBRH147 AF2042
GQ085455.1 gg_id:431512 Topographical and Temporal Human Skin Microbiome skin elbow clone nbw275h11 GQ085
EU133861.1 gg_id:249511 tallgrass prarie soil clone FFCH8183 EU1338
AY007722.1 gg_id:29482 Quadricoccus australiensis str. Ben 117; NCIMB 13738; CIP 107055 AY0077
AY212577.1 gg_id:99384 water 10 m downstream manure clone 126ds10 AY2125
EU015096.1 gg_id:236222 Investigation estrogen-degrading bioreactors membrane bioreactor suspension and EU0150
AB288886.1 gg_id:201491 termite gut homogenate clone BOf2-09 AB2888
AB487233.1 gg_id:562019 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNIR0 AB4872
FJ167461.1 gg_id:353414 microbiology granular sludge under autotrophic mixotrophic and heterotrophic denit FJ1674
AB237674.1 gg_id:140535 microorganisms deep subsurface groundwater sedimentary rock milieu clone HDBW AB2376
AJ582195.1 gg_id:93931 soil near uranium mill tailings clone Gitt-GS-125 AJ5821

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EU283471.1 gg_id:277656 Model Early Earth Harbors Deeply Divergent Divisions Anderson Lake clone A1227 EU2834
GQ979961.1 gg_id:589706 microbial biofilm soluble nitrate reducing bioreactor clone SRC_NRB032 GQ979
AB166784.1 gg_id:107623 EBPR sludge clone UTFS-OF08-18 AB1667
AB456210.1 gg_id:553408 Microorganisms involved anaerobic phenol degradation treatment synthetic coke-o AB4562
AM982545.1 gg_id:278568 microbial and slurry pig straw spent bedding clone PIST-AGC10 AM982
AB198538.1 gg_id:135858 termite gut wall fraction clone RsW02-065 AB1985
DQ088747.1 gg_id:131456 subsurface clone BE23FW031301A4RD-1 DQ088
FJ374248.1 gg_id:364266 Inter- and intraspecific humivorous scarab beetle larvae (Pachnoda spp.) larval hindg FJ3742
GQ105914.1 gg_id:447649 Topographical and Temporal Human Skin Microbiome skin gluteal crease clone nbw GQ105
DQ395985.1 gg_id:151035 deep-sea octacoral clone ctg_CGOGA28 DQ395
AJ005817.1 gg_id:6750 Thauera terpenica str. 58Eu AJ0058
AB192235.1 gg_id:141727 termite gut homogenate clone RsTu1-85 AB192
AJ863432.1 gg_id:120677 poplar tree microcosm rhizoplane flooded clone 23RPF22 AJ8634
GQ098414.1 gg_id:416580 Topographical and Temporal Human Skin Microbiome skin umbilicus clone nbw559c GQ098
AY188298.1 gg_id:80896 marine? clone KD2-110 AY1882
GQ104731.1 gg_id:415263 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ104
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AM268363.1 gg_id:185155 Functional Redundancy Versatility Rhodocyclales Activated Sludge: Improvement ar AM268
FJ439857.1 gg_id:513084 Investigation estrogen-degrading bioreactors membrane bioreactor biofilm clone M3 FJ4398
GQ103332.1 gg_id:463253 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw699b0 GQ103
FJ478630.1 gg_id:569356 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4786
GQ979958.1 gg_id:532793 microbial biofilm soluble nitrate reducing bioreactor clone SRC_NRB029 GQ979
AB166882.1 gg_id:104301 Azonexus sp. str. Slu-05 AB1668
GU454933.1 gg_id:520944 Understanding short-chain fatty acids accumulation enhanced alkaline digestion: kir GU454
AB192066.1 gg_id:143225 termite gut homogenate clone M1PL1-62 AB192
GQ065924.1 gg_id:397799 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw94e08 GQ065
GU454955.1 gg_id:525767 Understanding short-chain fatty acids accumulation enhanced alkaline digestion: kir GU454
DQ517100.1 gg_id:160794 Cultivation-intensive study ephemeral hypereutrophic aquatic microbial Mojave Des DQ517
GU222238.1 gg_id:586616 Anaerobic biodegradation branched hydrocarbon biomarker analog pristane wastew GU222
AY770497.1 gg_id:143561 and isolation that grow on 12-dichloroethane under nitrate reducing conditions sedi AY7704
AB486739.1 gg_id:585587 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSCOR AB4867
GQ231337.1 gg_id:560504 -stable isotope probing integrated metagenomics biphenyl dioxygenase genes polyc GQ231
AF011343.1 gg_id:6783 Azoarcus communis str. SWub3 AF0113
FJ823940.1 gg_id:529251 Controlling accumulation fermentation inhibitors biorefinery recycle water microbial FJ8239
AY662036.1 gg_id:136814 uranium-contaminated groundwater clone 300I-F02 AY6620
FJ479338.1 gg_id:539491 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4793
GQ108087.1 gg_id:430940 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ108
EU244087.1 gg_id:360080 16SrRNA probes river Leine sediment clone sl1373 EU2440
EF019494.1 gg_id:214657 aspen rhizosphere clone Elev_16S_763 EF0194
AB049763.1 gg_id:27927 Denitromonas aromaticus str. AS-7 AB0497

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AF047462.1 gg_id:6696 Dechloromonas agitata str. CKB AF0474
GQ389180.1 gg_id:576785 quality deterioration drinking system during serious red water outbreak clone D50 GQ389
GQ092099.1 gg_id:399670 Topographical and Temporal Human Skin Microbiome skin elbow clone nbw406a11c GQ092
EU104068.1 gg_id:321632 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
EU639302.1 gg_id:355190 Ecological Role Firmicutes Identified Thermophilic Microbial Fuel Cells thermophilic n EU6393
AM268352.1 gg_id:186954 Functional Redundancy Versatility Rhodocyclales Activated Sludge: Improvement ar AM268
GQ102268.1 gg_id:399299 Topographical and Temporal Human Skin Microbiome skin inguinal crease clone nbv GQ102
GQ074891.1 gg_id:413877 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ074
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GQ083179.1 gg_id:447534 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ083
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AB201043.1 gg_id:154231 Zoogloea oryzae str. -7 AB201
GQ104097.1 gg_id:424182 Topographical and Temporal Human Skin Microbiome skin glabella clone nbw563b0 GQ104
GQ093001.1 gg_id:451554 Topographical and Temporal Human Skin Microbiome skin elbow clone nbw407g11c GQ093
GQ103863.1 gg_id:381601 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ103
AM268341.1 gg_id:194016 Functional Redundancy Versatility Rhodocyclales Activated Sludge: Improvement ar AM268
EU834782.1 gg_id:591111 Microbiology Aerobic Phosphorus Removal Process aerobic phosphorus-removing se EU8347
AY689093.1 gg_id:103490 activated sludge clone KRZ70 AY6890
GQ099610.1 gg_id:377523 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw51 GQ099
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GQ104481.1 gg_id:381484 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ104
FJ375501.1 gg_id:540537 T-RFLP powerful tool rapid assessment cells microbial fuel cell sheet carbon paper an FJ3755
GQ102905.1 gg_id:421705 Topographical and Temporal Human Skin Microbiome skin inguinal crease clone nbv GQ102
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AF502233.1 gg_id:59335 EBPR sludge lab scale clone HP1A03 AF5022
GQ106704.1 gg_id:422891 Topographical and Temporal Human Skin Microbiome skin back clone nbw539e12c1 GQ106
GQ092947.1 gg_id:439576 Topographical and Temporal Human Skin Microbiome skin elbow clone nbw407b05c GQ092
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GQ095282.1 gg_id:374709 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ095
CU466885.1 gg_id:344216 Evry wastewater treatment plant anoxic basin clone CU466
D14254.1 gg_id:6731 Zoogloea ramigera str. IAM12136 D14254

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AM268353.1 gg_id:174268 Functional Redundancy Versatility Rhodocyclales Activated Sludge: Improvement ar AM268
AJ505851.1 gg_id:70483 Zoogloea resiniphila str. PIV-3A2w = DSM 14766 AJ5058
EF632935.1 gg_id:253630 Unique microbial contrasting aquatic altitude Andean Altiplano (northern Chile) fresh EF6329
EU133805.1 gg_id:249736 tallgrass prarie soil clone FFCH7204 EU1338
EF020008.1 gg_id:206405 aspen rhizosphere clone Elev_16S_1398 EF0200
EU589266.1 gg_id:266605 China Ningxia rice paddy field soil clone 1_E8 EU589
FJ748781.1 gg_id:567553 Vertical and sulfate-reducing prokaryotes estuarine Southern China Pearl River Estuai FJ7487
EU276554.1 gg_id:361100 -DGGE and Approaches Analyze Effect Agricultural Practices on Soil agricultural soil c EU2765
DQ499310.1 gg_id:159574 acidic cave snottite biofilm clone CV71 DQ499
FJ810598.1 gg_id:394416 Subsurface ecosystem resilience: Long-term attenuation subsurface contaminants su FJ8105
EF032745.1 gg_id:187854 cyanobacterial mat in lava cave clone HAVOmat01 EF0327
EF018556.1 gg_id:215521 aspen rhizosphere clone Amb_16S_819 EF0185
DQ351927.1 gg_id:165444 manganese oxidation soil clone JH-GY05 DQ351
EU287309.1 gg_id:254959 Pacific arctic surface sediment clone S26-9 EU2873
EU133892.1 gg_id:243583 tallgrass prarie soil clone FFCH4041 EU1338
FJ535105.1 gg_id:534686 microbial colonizing walls Karstic Slovenia cave wall biofilm clone 3PJM119 FJ5351
AB294327.1 gg_id:204721 Microbial structure deep coal seem groundwater stream clone SWB16 AB2943
GQ302538.1 gg_id:536176 prokaryotic sediment Shawan methods cold spring clone sw-xj248 GQ302
DQ413098.1 gg_id:153104 Microbial -dependent and Approaches Anaerobic/Aerobic SBR Reactor EBPR sludge DQ413
DQ228413.1 gg_id:144006 microbial compositions two subglacial possible role microbes chemical weathering p DQ228
AF280846.1 gg_id:17553 bioreactor clone mle1-7 AF2808
AB179499.1 gg_id:100324 sedimentary rock clone MIZ08 AB1794
GQ262859.1 gg_id:585913 Structure Simulated Level Waste Site simulated level waste site clone F1_34X GQ262
EU133840.1 gg_id:253509 tallgrass prarie soil clone FFCH1549 EU1338
AJ581593.1 gg_id:93464 soil near uranium mill tailings clone KCM-B-81 AJ5815
DQ906845.1 gg_id:242049 Oman subsurface soil clone 10A-2 DQ906
AM990454.1 gg_id:278532 Sterolibacterium sp. str. TKU1 AM990
AM935260.1 gg_id:543221 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM935
GQ261308.1 gg_id:552362 Study on Sediment sediment Dongping Lake clone 1S2-21 GQ261
GQ263997.1 gg_id:509604 Structure Simulated Level Waste Site simulated level waste site clone WC2_91 GQ263
AY555807.1 gg_id:102883 Bor Khlueng hot spring clone PK332 AY5558
EF516193.1 gg_id:219982 grassland soil clone FCPS563 EF5161
NR_024834.1 gg_id:573240 Thiobacter subterraneus str. C55; JCM 12421 NR_02
DQ413075.1 gg_id:150728 Microbial -dependent and Approaches Anaerobic/Aerobic SBR Reactor EBPR sludge DQ413
GQ263936.1 gg_id:593889 Structure Simulated Level Waste Site simulated level waste site clone WC2_188 GQ263
DQ906784.1 gg_id:251160 Oman subsurface soil clone 2B-6 DQ906
EF540388.1 gg_id:225930 oil-shale semi-coke solid waste clone MJK28 EF5403
EF018248.1 gg_id:214289 aspen rhizosphere clone Amb_16S_603 EF0182
GQ302536.1 gg_id:560421 prokaryotic sediment Shawan methods cold spring clone sw-xj169 GQ302
EF019282.1 gg_id:206997 aspen rhizosphere clone Elev_16S_468 EF0192
FJ478674.1 gg_id:510039 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4786

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EU881149.1 gg_id:352105 Karst region primary forest soil clone KPF200711-158 EU881
EU335200.1 gg_id:249281 Changes microbial metabolic and along hydrogeochemically variable profile saturate EU335
GQ263985.1 gg_id:545926 Structure Simulated Level Waste Site simulated level waste site clone WC2_75 GQ263
GQ302534.1 gg_id:540962 prokaryotic sediment Shawan methods cold spring clone sw-xj69 GQ302
DQ413095.1 gg_id:152559 Microbial -dependent and Approaches Anaerobic/Aerobic SBR Reactor EBPR sludge DQ413
DQ499286.1 gg_id:159638 acidic cave snottite biofilm clone CV30 DQ499
GQ263251.1 gg_id:585912 Structure Simulated Level Waste Site simulated level waste site clone F3_77X2 GQ263
DQ823210.1 gg_id:171073 Techniques Microbial Studies Oregon Caves National Monument Quad 17 on trail cl DQ823
EF019065.1 gg_id:214169 aspen rhizosphere clone Amb_16S_1641 EF019
FJ517093.1 gg_id:510574 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled water c FJ5170
EU133787.1 gg_id:257306 tallgrass prarie soil clone FFCH5275 EU133
EU449747.1 gg_id:575467 -DGGE and Approaches Analyze Effect Agricultural Practices on Soil agricultural soil c EU449
GQ262854.1 gg_id:593141 Structure Simulated Level Waste Site simulated level waste site clone F1_31X GQ262
GU208422.1 gg_id:588520 Study on Sediment Dongping Lake sediment clone Fh3-20 GU208
AY212696.1 gg_id:106348 water 10 m downstream manure clone 248ds10 AY212
DQ009366.1 gg_id:117907 bacterioplankton clone SPOTSAPR01_5m110 DQ009
EF019806.1 gg_id:205805 aspen rhizosphere clone Elev_16S_997 EF0198
AJ536880.1 gg_id:78574 uranium mining waste pile soil sample clone JG30-KF-AS41 AJ5368
EF516474.1 gg_id:220381 grassland soil clone FCPO565 EF5164
EU440652.1 gg_id:552054 Soil are Changed Agricultural Practices agricultural soil clone Plot17-F06 EU440
AY050578.1 gg_id:71612 MCB-contaminated groundwater-treating reactor clone GOUTA6 AY0505
EF492893.1 gg_id:243674 Iron-manganese Nodule and Its Surrounding Soil soil around iron-manganese nodule EF4928
EU753671.1 gg_id:341483 modern freshwater stromatolites Ruidera Pools Natural Park Spain dry stromatolite EU753
AJ519655.1 gg_id:76131 uranium mill tailings soil sample clone GuBH2-AD-29 proteobacterium AJ5196
AJ518781.1 gg_id:75238 uranium mining waste pile clone JG37-AG-7 proteobacterium AJ5187
EU133774.1 gg_id:244282 tallgrass prarie soil clone FFCH3050 EU133
DQ837228.1 gg_id:248208 Archaeal and pristine National Park Spain Donana coastal aquifer clone 49S1_1B_42 DQ837
EU133778.1 gg_id:246758 tallgrass prarie soil clone FFCH17581 EU133
AY922034.1 gg_id:114213 farm soil clone AKYG576 AY922
EF562187.1 gg_id:257648 Subsurface microbial deep-granitic-fracture water colorado Henderson Molybdenum EF5621
DQ499281.1 gg_id:159652 acidic cave snottite biofilm clone CV21 DQ499
FJ716013.1 gg_id:558969 Diet- gut microbiota Lumbricus rubellus fingerprinting technique and cloning intestina FJ7160
EF516456.1 gg_id:222917 grassland soil clone FCPS389 EF5164
AB179533.1 gg_id:101884 sedimentary rock clone MIZ42 AB179
DQ451461.1 gg_id:156973 forest soil clone FAC22 DQ451
GQ396825.1 gg_id:557255 Functional unvegetated perhumid recently-deglaciated soils do not correlate shifts s GQ396
EF019880.1 gg_id:204943 aspen rhizosphere clone Elev_16S_1242 EF0198
DQ404836.1 gg_id:154858 contaminated sediment clone 655965 DQ404
EU445220.1 gg_id:555446 Isolation Weight Forest Topsoil Metagenomic tropical forest topsoil clone R48 EU445
AY922053.1 gg_id:113596 farm soil clone AKYG676 AY922
DQ640725.1 gg_id:230327 full-scale EBPR sludge clone Skagen120 DQ640