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AM179906.1 gg_id:148234 Identification microflora (Oncorhynchus mykiss) rainbow trout intestine clone G3-1 AM179
GQ006791.1 gg_id:502872 Topographical and Temporal Human Skin Microbiome skin occiput clone nbu87g04c GQ006
AY947495.2 gg_id:114821 subgingival plaque clone OH1A AY947
AF439641.1 gg_id:35224 Veillonella atypica str. ATCC 17744 AF439
GQ027353.1 gg_id:421354 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ027
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GQ032280.1 gg_id:370470 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ032
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GQ084875.1 gg_id:431117 Topographical and Temporal Human Skin Microbiome skin manubrium clone nbw26 GQ084
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GQ114025.1 gg_id:385911 Topographical and Temporal Human Skin Microbiome skin retroauricular crease clo GQ114
AF439640.1 gg_id:30716 Veillonella parvula str. CIP 60.1 AF439
EF510428.1 gg_id:218682 Loss During Antibiotic Treatment 1 Intubated Patients Colonized Pseudomonas aeru EF5104
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GQ006604.1 gg_id:503059 Topographical and Temporal Human Skin Microbiome skin occiput clone nbu85e02c GQ006
GQ072552.1 gg_id:374904 Topographical and Temporal Human Skin Microbiome skin manubrium clone nbw19 GQ072
FJ557333.1 gg_id:518743 Survey Approach Assess Endotracheal Tube Biofilms and their Origins biofilm extubat FJ5573
FJ369852.1 gg_id:361643 feces adult twins and mothers; TS55 clone TS55_a01e08 FJ3698
FJ369825.1 gg_id:357386 feces adult twins and mothers; TS55 clone TS55_a01c04 FJ3698
FJ370053.1 gg_id:358239 feces adult twins and mothers; TS55 clone TS55_a03h07 FJ3700
AB478008.1 gg_id:576201 Microbial aquifer subsurface accretionary prism Japan geothermal groundwater clo AB478
AY538171.1 gg_id:98963 Desulfitobacter alkalitolerans str. Sk.kt5 AY538
FJ638550.1 gg_id:570010 Temperature dependent succession sulfate reducing muddy southwestern Taiwan h FJ6385
DQ230961.1 gg_id:142730 subsurface water clone DR938CH110701SACH36 DQ230
EU639075.1 gg_id:306415 Ecological Role Firmicutes Identified Thermophilic Microbial Fuel Cells thermophilic EU639
DQ413080.1 gg_id:153409 Microbial -dependent and Approaches Anaerobic/Aerobic SBR Reactor EBPR sludge DQ413
GQ502520.1 gg_id:542395 Effects Geographic Location on Formosan Subterranean Termite (FST) gut clone Cf4- GQ502
DQ988278.1 gg_id:192181 Microbial lab-scale A1-A2-O fixed biofilm system coking wastewater treatment with DQ988
EU638492.1 gg_id:319720 Ecological Role Firmicutes Identified Thermophilic Microbial Fuel Cells thermophilic EU638
AB118592.1 gg_id:93438 Endosymbiont 'TC1' Trimyema compressum clone TC1 AB118
FJ712502.1 gg_id:526485 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) FJ7125
AB439298.1 gg_id:541578 and microbial oils produced Asia petroleum crude oil (Daqing) imported China clone AB439
EU981271.1 gg_id:353812 sulfate reducing living bench scale aquifer column water clone E9_SW EU981
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X89072.1 gg_id:14311 Halocella cellulolsilytica str. DSM 7362 X89072
L22890.1 gg_id:14313 Halanaerobium salsuginis str. VS-752 L22890

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EF585982.1 gg_id:247996 Methanol-assimilating anaerobic solid waste digester fed methanol clone B55_K_B_A EF5859
EF558989.1 gg_id:244525 Microbial functional groups solid waste digestor stable isotope probing thermophilic EF5589
EU869420.1 gg_id:316746 Microbial hypersaline Sidi Ameur water Himalatt Salt Lake Algerian Sahara clone ARD EU8694
EU245629.1 gg_id:249184 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-M3-H0 EU2456
EF559026.1 gg_id:238840 Microbial functional groups solid waste digestor stable isotope probing thermophilic EF5590
EF106374.1 gg_id:201587 microbial two endoevaporitic salt crusts evaporitic crust Guerrero Negro Baja clone 2 EF1063
EU245738.1 gg_id:255254 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-M5-F07 EU2457
EF106367.1 gg_id:202585 microbial two endoevaporitic salt crusts evaporitic crust Guerrero Negro Baja clone 2 EF1063
FJ202891.1 gg_id:320426 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2028
AM947558.1 gg_id:264238 Minor changes detected microbial structure and after three week thermophilic dige AM947
X89077.1 gg_id:14308 Acetohalobium arabaticum X89077
NR_028804.1 gg_id:581342 Selenihalanaerobacter shriftii str. DSSE-1; ATCC BAA-73 NR_028
EF459724.1 gg_id:219578 Prokaryotic Hypersaline Inland Turkey Tuz Lake (hypersaline) clone 4-62B EF4597
EU327343.1 gg_id:288770 Halobacteroides sp. str. 6SANG EU3273
X89074.1 gg_id:14301 Halobacteroides halobius str. DSM 5150 X89074
GU452712.1 gg_id:525021 Natroniella sp. str. AHT18 GU452
X89073.1 gg_id:14302 Orenia marismortui str. DSM 5156 X89073
Y18485.1 gg_id:69790 Orenia salinaria str. SG3902 Y18485
EU245188.1 gg_id:242318 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-H3-G09 EU2451
EF559012.1 gg_id:250397 Microbial functional groups solid waste digestor stable isotope probing thermophilic EF5590
AM947554.1 gg_id:270282 Minor changes detected microbial structure and after three week thermophilic dige AM947
EF586067.1 gg_id:246373 Methanol-assimilating anaerobic solid waste digester fed methanol clone M55_D21_ EF5860
EF558957.1 gg_id:238729 Microbial functional groups solid waste digestor stable isotope probing thermophilic EF5589
EF586053.1 gg_id:257232 Methanol-assimilating anaerobic solid waste digester fed methanol clone M55_D21_ EF5860
EU878323.1 gg_id:567033 Investigation factors influencing production large-scale municipal biogas plant therm EU8783
EU639242.1 gg_id:313846 Ecological Role Firmicutes Identified Thermophilic Microbial Fuel Cells thermophilic n EU6392
FN436027.1 gg_id:539789 Molekularbiologische Charakterisierung der bakteriellen Populationsdynamik eines t FN4360
AM947555.1 gg_id:266221 Minor changes detected microbial structure and after three week thermophilic dige AM947
EF559059.1 gg_id:243394 Microbial functional groups solid waste digestor stable isotope probing thermophilic EF5590
EF586029.1 gg_id:252533 Methanol-assimilating anaerobic solid waste digester fed methanol clone M35_D20_ EF5860
EF559123.1 gg_id:226234 mesophilic anaerobic digester clone C35_D63_H_B_F12 EF5591
EF559004.1 gg_id:248896 Microbial functional groups solid waste digestor stable isotope probing thermophilic EF5590
EF586046.1 gg_id:241887 Methanol-assimilating anaerobic solid waste digester fed methanol clone M55_D15_ EF5860
GU455299.1 gg_id:522555 Understanding short-chain fatty acids accumulation enhanced alkaline digestion: kir GU455
EU358734.1 gg_id:312486 structure archaeal denaturing gel electrophoresis and sequencing mesophilic biogas EU3587
EF422412.2 gg_id:224884 Dethiobacter alkaliphilus str. AHT1 EF4224
AF507887.1 gg_id:62675 Mono Lake at depth 35m station 6 July 2000 clone ML635J-4 AF5078
AJ295640.1 gg_id:31598 uranium mill tailing clone KF-Gitt2-16 AJ2956
AJ582199.1 gg_id:93440 soil near uranium mill tailings clone Gitt-GS-113 AJ5821
DQ432150.1 gg_id:156668 prokaryotic alkaline hypersaline lakes Wadi Natrun Egypt Lake Fazda sediment clone DQ432
AF507888.1 gg_id:55772 Mono Lake at depth 35m station 6 July 2000 clone ML635J-65 G+C AF5078

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FJ203016.1 gg_id:351673 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2030
GQ848219.1 gg_id:541684 and Archaeal Xiarinur Soda Lake Water-sediment Interface water-sediment interfac GQ848
EU812988.1 gg_id:341763 Microbial structure thermophilic reactors biogas reactor fed corn silage clone TUM-N EU8129
EU592468.1 gg_id:573525 Seasonal changes Salton Sea hypersaline sediment clone SSS47A EU5924
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GQ848204.1 gg_id:588856 and Archaeal Xiarinur Soda Lake Water-sediment Interface water-sediment interfac GQ848
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AJ271451.1 gg_id:73970 Natronoanaerobium halophilum G-M14CH-4 AJ2714
AJ271450.1 gg_id:23520 Natronoanaerobium salstagnum O-M12SP-2 AJ2714
AJ271452.1 gg_id:72206 Natronoanaerobium aggerbacterium G-M16NWC-4 AJ2714
DQ354747.1 gg_id:155876 subsurface water clone MS149BH1062003_30 DQ354
AB539949.1 gg_id:571087 Evidence syntrophic acetate oxidation coupled hydrogenotrophic methanogenesis -t AB5399
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DQ125604.1 gg_id:134479 uranium contaminated soil clone AKAU3609 DQ125
GQ402678.1 gg_id:512997 Accessing microbial biodiversity and metabolic potential tropical -based approach p GQ402
DQ125579.1 gg_id:132342 uranium contaminated soil clone AKAU3574 DQ125
DQ125547.1 gg_id:129508 uranium contaminated soil clone AKAU3523 DQ125
DQ887916.1 gg_id:168335 thermophilic anaerobic solid waste bioreactor clone GHB3B06 DQ887
DQ125708.1 gg_id:133296 uranium contaminated soil clone AKAU3802 DQ125
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EF585978.1 gg_id:246509 Methanol-assimilating anaerobic solid waste digester fed methanol clone B55_K_B_A EF5859
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DQ125736.1 gg_id:134788 uranium contaminated soil clone AKAU3845 DQ125
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CU919791.1 gg_id:534056 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU919
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CU919182.1 gg_id:566892 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU919

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AY684090.1 gg_id:130626 methanogenic enrichment 'MRE50' clone MRE50b08 AY684
CT573824.1 gg_id:150758 Evry municipal wastewater treatment plant clone 013H11_B_SD_P15 CT573
AY526502.1 gg_id:136583 Microbial thermophilic anaerobic sludge methanol-fed lab scale bioreactor clone D3 AY526
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CU918907.1 gg_id:539546 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU918
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EF198048.1 gg_id:185617 Microbial structures mesophilic and methanogenic consortia anaerobic thermophilic EF198
EU639113.1 gg_id:306099 Ecological Role Firmicutes Identified Thermophilic Microbial Fuel Cells thermophilic n EU639
AB374111.1 gg_id:246025 Effect temperature on microbial glucose-degrading methanogenic consortium under AB374
CR933199.1 gg_id:110387 Evry municipal wastewater treatment plant clone 008A06_B_SD_P15 CR933
CU919263.1 gg_id:549429 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU919
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AB234270.1 gg_id:145732 Biotransformation Polychlorinated Dioxins Microbial Dynamics Sediment Microcosm AB234
DQ018815.1 gg_id:132872 Unq Mcrbl Anrbc Swn Lgns anrbc swn lgn cln C-146 DQ018
EF586045.1 gg_id:245235 Methanol-assimilating anaerobic solid waste digester fed methanol clone M55_D15_ EF586
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AJ004811.1 gg_id:16095 Caldicellulosiruptor kristjanssonii str. I77R1B AJ0048
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FJ189575.1 gg_id:351375 digester anaerobic sludge clone EUB_54 FJ1895

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DQ152922.1 gg_id:139369 Caldicellulosiruptor sp. str. YI5 DQ152
CP001785.1 gg_id:521670 Ammonifex degensii str. KC4 CP0017
EF554597.1 gg_id:233108 Fervidomicrobium thiophilum str. SR EF5545
NC_003869.1 gg_id:471376 Thermoanaerobacter tengcongensis MB4 str. MB4T NC_00
AF174484.1 gg_id:39044 Carboxydibrachium pacificum str. JM AF1744
EU109461.1 gg_id:237130 Thermoanaerobacter subterraneus str. S2302 EU1094
AF212925.1 gg_id:26976 Thermoanaerobacter yonseiensis str. KB1 AF2129
EF554599.1 gg_id:234626 Thermoanaerobacter subterraneus str. 2707 EF5545
EF195126.1 gg_id:175440 Caldanaerobacter hydrothermalis str. K67 EF1951
FJ189534.1 gg_id:349650 digester anaerobic sludge clone EUB_13 FJ1895
EF554598.1 gg_id:232884 Thermoanaerobacter subterraneus str. SP47 EF5545
AY278483.1 gg_id:87364 Thermoanaerobacter keratinophilus str. 2KXI AY2784
DQ115806.1 gg_id:134076 Thermoanaerobacter tengcongensis str. 185y2 DQ115
CP000141.1 gg_id:140847 Carboxydothermus hydrogenoformans str. Z-2901 CP0001
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EU638454.1 gg_id:309145 Ecological Role Firmicutes Identified Thermophilic Microbial Fuel Cells thermophilic n EU6384
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DQ647145.1 gg_id:257640 Microbial structure -temperature oil-field produced water Troll formations North Se DQ647
EF558950.1 gg_id:246613 Microbial functional groups solid waste digestor stable isotope probing thermophilic EF5589
EU386162.1 gg_id:265020 Thermacetogenium sp. str. Sp3 EU3861
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AY701757.1 gg_id:106184 Thermoanaerobacter sp. str. AIP 504.99 AY7017
NC_010321.1 gg_id:259585 Thermoanaerobacter pseudethanolicus 33223 str. ATCC 33223; 39E NC_010
EU262599.1 gg_id:240738 Thermoanaerobacter inferii str. AK15 EU2625
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DQ681066.1 gg_id:163311 Thermoanaerobacter wiegelii str. B5 DQ681
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EU731003.1 gg_id:338174 single-species ecosystem deep within Earth fracture water borehole MP104 level 10 EU731
FJ712408.1 gg_id:523660 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) k FJ7124
EU730994.1 gg_id:353154 single-species ecosystem deep within Earth fracture water borehole MP104 level 10 EU730
EU419136.1 gg_id:267107 Microbial and ecology selected andesitic hydrothermal acidic hydrothermally modifi EU419
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AB062280.1 gg_id:20457 Thermanaeromonas toyohensis str. ToBE AB062
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EU638335.1 gg_id:348663 Ecological Role Firmicutes Identified Thermophilic Microbial Fuel Cells thermophilic n EU638
FJ638536.1 gg_id:572985 Temperature dependent succession sulfate reducing muddy southwestern Taiwan hc FJ6385
AB374120.1 gg_id:243918 Effect temperature on microbial glucose-degrading methanogenic consortium under AB374
AB434897.1 gg_id:271632 Effect temperature on microbial glucose-degrading methanogenic consortium under AB434
AE013109.1 gg_id:66678 Thermoanaerobacter tengcongensis str. MB4 AE013
AB374132.1 gg_id:243488 Effect temperature on microbial glucose-degrading methanogenic consortium under AB374
AB374117.1 gg_id:238691 Effect temperature on microbial glucose-degrading methanogenic consortium under AB374
AB374134.1 gg_id:256314 Effect temperature on microbial glucose-degrading methanogenic consortium under AB374
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EU639339.1 gg_id:323523 Ecological Role Firmicutes Identified Thermophilic Microbial Fuel Cells thermophilic n EU639
EF174273.1 gg_id:190840 direct activated sludge compost maturation stage clone H42-814 EF1742
CU920710.1 gg_id:539239 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU920
DQ079649.1 gg_id:137457 deep terrestrial subsurface fluid-filled fracture clone EV818FW062101BH4MD9 DQ079
AB260059.1 gg_id:179484 Microbial black exposed hot ridge flank crustal fluids Black rust formation on boreho AB260
AB300079.1 gg_id:220063 Wide ancient microbial DNAs Quaternary sediments holocene mud sediment clone A AB300
AM777947.1 gg_id:237589 nonsaline . Influence pore-size filters determination structure alkaline groundwater AM777
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DQ088772.1 gg_id:131486 subsurface clone BE325FW032701CTS_hole1-1 DQ088
AB426224.1 gg_id:275227 -situ identification study benzene-degrading s under methanogenic condition anaerc AB426
EF632928.1 gg_id:246403 Unique microbial contrasting aquatic altitude Andean Altiplano (northern Chile) fresh EF6329
FJ203601.1 gg_id:339359 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2036
AB539950.1 gg_id:517375 Evidence syntrophic acetate oxidation coupled hydrogenotrophic methanogenesis -t AB539
DQ256302.1 gg_id:142791 subsurface water clone EV818BHEB6102702SAFF42 DQ256

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EF585979.1 gg_id:257911 Methanol-assimilating anaerobic solid waste digester fed methanol clone B55_K_B_/ EF5859
EF559068.1 gg_id:252254 Microbial functional groups solid waste digestor stable isotope probing thermophilic EF5590
AM947532.1 gg_id:265589 Minor changes detected microbial structure and after three week thermophilic dige AM947
AF282252.1 gg_id:17524 Ferribacter thermoautotrophicus str. JW/JH-Fiji-2 AF2822
DQ815026.1 gg_id:169250 zebrafish digestive tract clone aab49a06 DQ815
DQ815290.1 gg_id:171837 zebrafish digestive tract clone aab52g10 DQ815
DQ815286.1 gg_id:171805 zebrafish digestive tract clone aab52g04 DQ815
DQ815110.1 gg_id:173268 zebrafish digestive tract clone aab50b10 DQ815
DQ815013.1 gg_id:161980 zebrafish digestive tract clone aab56g09 DQ815
AM179923.1 gg_id:145422 Identification microflora (Oncorhynchus mykiss) rainbow trout intestine clone G15-: AM179
DQ815269.1 gg_id:172292 zebrafish digestive tract clone aab52e06 DQ815
DQ815170.1 gg_id:168343 zebrafish digestive tract clone aab51a08 DQ815
DQ815292.1 gg_id:170458 zebrafish digestive tract clone aab52h02 DQ815
EU460633.1 gg_id:297393 polar bear feces clone PB2_aai23f09 EU460
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EU460631.1 gg_id:298501 polar bear feces clone PB2_aai23f07 EU460
EU460679.1 gg_id:297037 polar bear feces clone PB2_aai24c12 EU460
EU460629.1 gg_id:293351 polar bear feces clone PB2_aai23f05 EU460
GU293194.1 gg_id:562985 Microbial contents and mucus yel catfish (Pelteobagrus fulvidraco) stomach content GU293
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EU460361.1 gg_id:297672 polar bear feces clone PB1_aai27c07 EU460
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NR_025679.1 gg_id:554906 Fusobacterium canifelinum str. RMA 12708; CCUG 49735 Q11 NR_02
AJ295750.1 gg_id:34266 Fusobacterium equinum str. horse VPB 4027 AJ2957
X74408.1 gg_id:11309 Fusobacterium necrophorum str. B FnS-40 X74408
X74407.1 gg_id:11305 Fusobacterium necrophorum str. AB FnS-1 X74407
FJ471661.1 gg_id:524898 Fusobacterium nucleatum str. ChDC F319 FJ4716
GQ004485.1 gg_id:505177 Topographical and Temporal Human Skin Microbiome skin nare clone nbu59b02c1 GQ004
EU772809.1 gg_id:306442 Bush dog feces clone bdog1_aai77h07 EU7728
AF543300.1 gg_id:58280 Fusobacterium nucleatum str. ChDC OS50 AF5433
GQ448468.1 gg_id:511044 Microbiome neonatal calves: shedding microorganisms fecal clone calf784_2wks_gr GQ448
AF481217.1 gg_id:68347 Fusobacterium nucleatum str. E9_12 AF4812

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NZ_ACET01000043.1 gg_id:470433 Fusobacterium gonidiaformans str. ATCC 25563 NZ_AC
FJ471649.1 gg_id:529903 Fusobacterium nucleatum str. ChDC F290 FJ4716
GQ091870.1 gg_id:449592 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ091
GQ113263.1 gg_id:375228 Topographical and Temporal Human Skin Microbiome skin retroauricular crease clor GQ113
FJ978721.1 gg_id:408384 feces clone CL_F_417 FJ9787
EU774026.1 gg_id:308389 Cheetah feces clone CE3_aai08d02 EU774
EF089177.1 gg_id:185041 Fusobacterium nucleatum str. Ulm 14 EF0891
FJ978724.1 gg_id:381867 feces clone CL_F_422 FJ9787
FJ978541.1 gg_id:427042 feces clone CL_F_080 FJ9785
FJ471639.2 gg_id:514347 Fusobacterium nucleatum str. ChDC F113 FJ4716
GQ088072.1 gg_id:459200 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ088
FJ471673.1 gg_id:520375 Fusobacterium periodonticum str. ChDC F334 FJ4716
DQ440563.1 gg_id:156573 Fusobacterium nucleatum str. Ulm DQ440
GQ087757.1 gg_id:395408 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ087
GQ014795.1 gg_id:494871 Topographical and Temporal Human Skin Microbiome skin palm clone nbw764c09c1 GQ014
FJ978525.1 gg_id:468118 feces clone CL_F_056 FJ9785
CP001685.1 gg_id:529857 Leptotrichia buccalis C-1013-b str. DSM 1135 CP001
FJ577251.2 gg_id:568972 Leptotrichia hofstadii str. F0254 FJ5772
AF206305.1 gg_id:71541 Leptotrichia trevisanii AF206
AY134896.1 gg_id:59574 oral periodontitis clone EX103 AY1348
AM420035.1 gg_id:241454 Noma subgingival plaque clone 201G10(oral) AM420
CP001685.1 gg_id:529859 Leptotrichia buccalis C-1013-b str. DSM 1135 CP001
GQ129984.1 gg_id:399980 Genetic inventory 'JPL-SAF' during MSL mission spacecraft assembly clean room floc GQ129
AY207053.1 gg_id:83573 human mouth clone P2PB_51 AY207
AF432138.1 gg_id:31235 tongue dorsum scrapings clone FP036 AF432
GQ000682.1 gg_id:469247 Topographical and Temporal Human Skin Microbiome skin interdigital web space clc GQ000
GQ000882.1 gg_id:469447 Topographical and Temporal Human Skin Microbiome skin interdigital web space clc GQ000
AY349385.1 gg_id:92131 human oral cavity clone GT020 AY349
FJ976353.1 gg_id:578509 Interindividual healthy human oral microbiota mouth clone 12BB57 FJ9763
AM420193.1 gg_id:258023 Noma subgingival plaque clone 502D06(oral) AM420
GQ016176.1 gg_id:493491 Topographical and Temporal Human Skin Microbiome skin gluteal crease clone nbw GQ016
FJ577259.2 gg_id:566658 Leptotrichia goodfellowii str. F0264 FJ5772
AY029803.1 gg_id:70629 Leptotrichia hofstadii str. LB23 AY0298
AF287814.1 gg_id:38061 human subgingival plaque clone F138 AF2878
GQ032449.1 gg_id:403062 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ032
AF385571.1 gg_id:70038 crevicular epithelial cells clone EI013 AF3855
GQ016227.1 gg_id:493440 Topographical and Temporal Human Skin Microbiome skin gluteal crease clone nbw GQ016
EU137409.1 gg_id:267538 Bartonella-positive fleas: and assembly patterns Oropsylla hirsuta (prairie dog flea) C EU1374
NZ_ACVB02000014.1 gg_id:484462 Leptotrichia hofstadii str. F0254 NZ_AC
AM420110.1 gg_id:256867 Noma subgingival plaque clone 303F08(oral) AM420

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GQ422731.1 gg_id:555711 Leptotrichia sp. oral taxon 225 str. FEA2 GQ422
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GU086183.1 gg_id:593385 Leptotrichia shahii str. PW1036 GU086
NR_026243.1 gg_id:557791 Propionigenium maris str. 10succ1; DSM 9537 NR_02
EU236316.1 gg_id:296908 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg1bD2 EU2363
X54275.1 gg_id:11301 Propionigenium modestum X54275
AJ877255.1 gg_id:112138 Ilyobacter psychrophilus str. FQ50 AJ8772
AJ307980.1 gg_id:52873 Ilyobacter insuetus str. DSM 6831 T AJ3079
Y16799.1 gg_id:42042 Propionigenium maris str. DSL-1 Y16799
FJ264772.1 gg_id:324952 Manganese and Iron Dependent Anaerobic Oxidation Methane methane seep sedime FJ2647
FJ717186.1 gg_id:510621 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United Kingc FJ7171
FJ717188.1 gg_id:558150 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United Kingc FJ7171
EU052253.1 gg_id:240997 degradation kinetics and current efficiency cells supplied anode microbial fuel cell fe EU052
AY770718.1 gg_id:110394 Fusobacteria sea urchin Paracentrotus lividus clone PlS49 AY7707
CP001739.1 gg_id:518403 Sebaldella termitidis str. ATCC 33386 CP0017
EF608534.1 gg_id:228694 predatory Poecilus chalcites their response lab rearing and antibiotic treatment diges EF6085
EF218611.1 gg_id:212738 Leptotrichia amnionii str. CCUG 51847 EF2186
AY078425.1 gg_id:29566 Leptotrichia amnionii str. AMN-1 AY0784
AJ344093.1 gg_id:58542 Sneathia sanguinegens str. CCUG 41628T AJ3440
AB330759.1 gg_id:236558 Streptobacillus moniliformis str. KWG2 AB3307
GQ072215.1 gg_id:447942 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw193e1 GQ072
GQ850557.1 gg_id:591820 surface water northern Bering Sea clone u114 GQ850
FJ456795.1 gg_id:545014 microbiota two Antarctic notothenioid fish species intestinal flora clone A007_NCI FJ4567
DQ206413.1 gg_id:139839 microbial mediates arsenate reduction soda lake water clone ML-AsS-15 DQ206
EU044321.1 gg_id:256629 uences structure fosters richness subtropical Altamaha Ohopee River Valley inland d EU0443
EF019213.1 gg_id:213629 aspen rhizosphere clone Amb_16S_1909 EF0192
DQ490008.1 gg_id:158907 Microbial Great Boiling Spring sediment hot spring; temperature 80 C clone GBS_L1_ DQ490
EU134698.1 gg_id:255244 tallgrass prarie soil clone FFCH14117 EU1346
FM873495.1 gg_id:363705 occupant source house floor dust clone FD04E09 FM873
EF018217.1 gg_id:216704 aspen rhizosphere clone Amb_16S_564 EF0182
EF540422.1 gg_id:220442 oil-shale semi-coke solid waste clone MK51 EF5404
GQ113134.1 gg_id:433730 Topographical and Temporal Human Skin Microbiome skin glabella clone nbw675f06 GQ113
EU283562.1 gg_id:278836 Model Early Earth Harbors Deeply Divergent Divisions Anderson Lake clone A1853 EU2835
FJ444672.1 gg_id:591271 Cotton Rhizosphere cotton rhizosphere clone 2y-23 FJ4446
AB240260.1 gg_id:142399 microbial structure rhizosphere Phragmites bulk soil reed bed reactor laboratory clo AB2402
DQ640655.1 gg_id:234041 full-scale EBPR sludge clone Skagenf3 DQ640
GQ090575.1 gg_id:440096 Topographical and Temporal Human Skin Microbiome skin popliteal fossa clone nbw GQ090
AB234279.1 gg_id:146305 Biotransformation Polychlorinated Dioxins Microbial Dynamics Sediment Microcosm AB2342

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EF157193.1 gg_id:243247 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF1571
EF018812.1 gg_id:214676 aspen rhizosphere clone Amb_16S_1283 EF0188
FJ478758.1 gg_id:512313 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4787
EF203205.1 gg_id:207957 phylotypes sediments Lake Kastoria Greece sediment clone Kas174B EF2032
AF432635.1 gg_id:99232 lodgepole pine rhizosphere soil British Columbia Ministry Forests Long-Term Soil Prod AF4326
DQ067025.1 gg_id:133200 Lake Washington sediment clone pLW-82 DQ067
FJ536876.1 gg_id:587747 Linking performance microbiology biofilters treating dimethyl sulphide presence and FJ5368
DQ520169.1 gg_id:160270 shal China sequences hypertrophic freshwater lake clone ML-5-110.2 DQ520
EF018312.1 gg_id:207648 aspen rhizosphere clone Amb_16S_675 EF0183
AF545649.1 gg_id:72899 shortgrass steppe soil (Stunner series fine loamy over sandy Ustic Calciargid) under Bc AF5456
AY218696.1 gg_id:81623 penguin droppings sediments clone KD8-87 AY2186
AB426207.1 gg_id:264973 -situ identification study benzene-degrading s under methanogenic condition anaerc AB4262
EU979110.1 gg_id:336425 Effects intercropping Rhizobium inoculation on microbial biomass and rhizosphere b EU9791
DQ906809.1 gg_id:239203 Oman subsurface soil clone 5B-5 DQ906
EF516323.1 gg_id:225306 grassland soil clone FCPT668 EF5163
EF019647.1 gg_id:215970 aspen rhizosphere clone Elev_16S_798 EF0196
DQ067020.1 gg_id:129424 Lake Washington sediment clone pLW-79 DQ067
FJ479207.1 gg_id:588566 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4792
EF019081.1 gg_id:204271 aspen rhizosphere clone Amb_16S_1670 EF0190
FJ152750.1 gg_id:339480 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX5A_ FJ1527
EU044086.1 gg_id:237353 fluences structure fosters richness subtropical Altamaha Ohopee River Valley inland EU0440
EU193066.1 gg_id:360943 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU1930
AM934748.1 gg_id:549433 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM934
AY921939.1 gg_id:114406 farm soil clone AKYG583 AY9219
EU881263.1 gg_id:329628 (long-term grazing and burned annual) Karst region serious disturbance field soil clor EU8812
AY922055.1 gg_id:114310 farm soil clone AKYG703 AY9220
AY921712.1 gg_id:112987 farm soil clone AKYG963 AY9217
AY921894.1 gg_id:114101 farm soil clone AKYG1555 AY9218
EF612383.1 gg_id:237047 historic semiarid lead-zinc mine tailings site soil clone OS-C17 EF6123
AY395378.1 gg_id:103904 pasture soil clone EB1059 AY3953
AY192277.1 gg_id:79951 Rocky Mountain alpine soil clone W2a-1F AY1922
GU359099.1 gg_id:518957 rhizosphere soil peanut clone 2G11 GU359
DQ906907.1 gg_id:253096 Oman subsurface soil clone 12B-7 DQ906
EU801865.1 gg_id:314599 It's all ranking aquatic Chesapeake Bay MD clone 3C003242 EU8018
FJ746162.1 gg_id:512463 Microbial oligotrophic marine sediments: South Pacific Gyre ocean sediment 5306 m FJ7461
EF020040.1 gg_id:211950 aspen rhizosphere clone Elev_16S_1437 EF0200
FJ230809.1 gg_id:325203 Microclimate and limits photosynthesis cyanobacteria Northern Australia hypolithic s FJ2308
AB257644.1 gg_id:259192 Microbial few millimeters be surface dolomite rock 'Val Piora' Alps 1965 meters abo AB2576
AY921918.1 gg_id:113407 farm soil clone AKYG1585 AY9219
EU283563.1 gg_id:279650 Model Early Earth Harbors Deeply Divergent Divisions Anderson Lake clone A1336 EU2835

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EU652636.1 gg_id:277713 Seasonal variation microbial Yel Sea sediment clone D8S-116 EU6526
EU048610.1 gg_id:254529 Microbial surface marine sediment Xisha Trough South China Sea clone MD2902-B3 EU0486
AY922130.1 gg_id:114403 farm soil clone AKYH466 AY9221
AY673312.1 gg_id:110882 Gemmatimonadetes Ellin7146 AY6733
FJ479068.1 gg_id:560811 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4790
AY922123.1 gg_id:113603 farm soil clone AKYH1517 AY9221
EF020045.1 gg_id:207989 aspen rhizosphere clone Elev_16S_1446 EF0200
AY921994.1 gg_id:112986 farm soil clone AKYG1838 AY9219
GQ396903.1 gg_id:557467 Functional unvegetated perhumid recently-deglaciated soils do not correlate shifts s GQ396
AY921912.1 gg_id:114146 farm soil clone AKYG1614 AY9219
EU676443.1 gg_id:289924 rhizosphere pioneer plants (Bahia xylopoda and Viguiera linearis) growing on metals- EU6764
EF018145.1 gg_id:208048 aspen rhizosphere clone Amb_16S_479 EF0181
GU205280.1 gg_id:516178 unusual Microbial Venezuelan Tepui (Orthoquartzite) Cave: Dominance Chloroflexal GU205
FJ152768.1 gg_id:335807 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX5A_ FJ1527
AY289480.1 gg_id:90194 soil clone ABS-650 AY2894
GQ113661.1 gg_id:384470 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ113
EF516935.1 gg_id:219517 grassland soil clone FCPN548 EF5169
EU193059.1 gg_id:360637 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU1930
DQ532275.1 gg_id:168856 Geographical Variation Microbial Burden and Spacecraft Assembly Johnson Space C DQ532
EF688378.1 gg_id:250371 soil clone F6-84 EF6883
AY921776.1 gg_id:113308 farm soil clone AKYG961 AY9217
EU234231.1 gg_id:242383 river and un techniques downstream Wang Yang River receiving penicillin G producti EU2342
EU881123.1 gg_id:308062 Karst region primary forest soil clone KPF200711-086 EU8811
DQ067019.1 gg_id:128235 Lake Washington sediment clone pLW-78 DQ067
EU134880.1 gg_id:249280 tallgrass prarie soil clone FFCH2651 EU1348
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AY921771.1 gg_id:113042 farm soil clone AKYG666 AY9217
EF492900.1 gg_id:246196 Iron-manganese Nodule and Its Surrounding Soil soil around iron-manganese nodule EF4929
EU134828.1 gg_id:249948 tallgrass prarie soil clone FFCH6230 EU1348
EU861901.1 gg_id:325238 effects chronic fertilization on alpine tundra microbial implications carbon and cyclin EU8619
FJ755752.1 gg_id:565312 vertical archaeal water and Lake Taihu sediment clone LT-SB-B6 FJ7557
AY218678.1 gg_id:82251 penguin droppings sediments clone KD8-105 AY2186
AY921899.1 gg_id:114389 farm soil clone AKYG781 AY9218
DQ906808.1 gg_id:243477 Oman subsurface soil clone 5B-4 DQ906
GQ263232.1 gg_id:590444 Structure Simulated Level Waste Site simulated level waste site clone F3_42X GQ263
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GQ263030.1 gg_id:576743 Structure Simulated Level Waste Site simulated level waste site clone F2_153X GQ263
EF174264.1 gg_id:197816 direct activated 4th maturation stage sludge clone F37-814 EF1742

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GQ262961.1 gg_id:538070 Structure Simulated Level Waste Site simulated level waste site clone F1_92X GQ262
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GQ346931.1 gg_id:591890 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ346
DQ295242.1 gg_id:148154 Fragments Six Marine Planktonic Antarctic sea water clone Ant4D5 DQ295
AJ567595.1 gg_id:89477 deep-sea sediment clone MBAE42 AJ5675
EU287180.1 gg_id:251953 Pacific arctic surface sediment clone P13-87 EU287
EU374078.1 gg_id:276624 southern Cretan margin (eastern Mediterranean Sea) canyon and slope sediment clc EU374
AM997489.1 gg_id:548429 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
FJ205357.1 gg_id:512804 Prokaryotic deep-sea hydrothermal region East Lau Spreading Centre deep marine se FJ2053
AM935228.1 gg_id:574770 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM935
EU652634.1 gg_id:277772 Seasonal variation microbial Yel Sea sediment clone A8S-129 EU652
FJ746291.1 gg_id:572935 Microbial oligotrophic marine sediments: South Pacific Gyre ocean sediment 5306 m FJ7462
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EU287420.1 gg_id:244505 Pacific arctic surface sediment clone S26-120 EU287
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AB294923.1 gg_id:226129 -dependent & analyses hydrothermal system off Taketomi Island Japan microbial ma AB294
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AY493976.1 gg_id:136870 Fingerprinting aggregates soil clone 5 AY4939
DQ431899.1 gg_id:208663 microorganisms catalyzing nitrogen removal and organic matter metabolism perme DQ431
EU491861.1 gg_id:270789 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU4918

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EU289488.1 gg_id:254895 metagenomic approach biosynthesis ansamycin endophytes or symbionts enriched s EU2894
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EU652631.1 gg_id:277923 Seasonal variation microbial Yel Sea sediment clone C8S-21 EU6526
EU245438.1 gg_id:237565 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-M1-F03 EU2454
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EU219931.1 gg_id:247892 study prokaryotic approach landfill soil 5.5 ft. depth clone C6B EU2199
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AB491850.1 gg_id:542631 affiliation (Litopenaeus vannamei) Banyuwangi East Java microbial flocs floating brac AB4918
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EU134863.1 gg_id:255390 tallgrass prarie soil clone FFCH17465 EU1348
GU056071.1 gg_id:579069 Biologic Techniques Applied Microbial Prospecting China: Ban 876 Gas Oil Field Bein GU056
FJ152883.1 gg_id:351400 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX4CI FJ1528
AB250557.1 gg_id:160916 and ecology Grotta Azzurra Palinuro Cape (Salerno Italy) microbial mat clone #2-2 AB2505
EU283598.1 gg_id:279155 Model Early Earth Harbors Deeply Divergent Divisions Anderson Lake clone A406 EU2835
FJ205371.1 gg_id:510313 Prokaryotic deep-sea hydrothermal region East Lau Spreading Centre deep marine se FJ2053
EU700157.1 gg_id:288295 Study on Shantou Bay Marine Sediments Microorganisms sediment clone STU15 EU7001

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GQ347441.1 gg_id:581615 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
DQ234160.2 gg_id:144617 determined library mangrove clone DS076 DQ234
DQ329577.1 gg_id:159133 Guerrero Negro hypersaline microbial mat clone 05D2Z07 DQ329
DQ397339.1 gg_id:159241 Guerrero Negro hypersaline microbial mat clone 02D2Z08 DQ397
AB192119.1 gg_id:142691 termite gut homogenate clone M2PT2-27 AB192
FJ791860.1 gg_id:571248 Unexpectedly archaeal species shift between rare and dominant over thousand year FJ7918
AY197400.1 gg_id:86409 Guaymas Basin hydrothermal vent sediments clone B02R018 AY197
FJ959985.1 gg_id:579816 bottlenose dolphin upper respiratory tract blowhole clone D940_045 FJ9599
AB232534.1 gg_id:154098 Prokaryotic gills deep-sea hydrothermal vent shrimp gill Alvinocaris longirostris clone AB232
EU052256.1 gg_id:256207 degradation kinetics and current efficiency cells supplied anode microbial fuel cell fe EU052
AF367482.1 gg_id:17620 deep-sea hydrothermal vent clone VC1.2-cl02 AF367
GQ349327.1 gg_id:562277 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ349
EU265935.1 gg_id:537633 transition and zone within near-shore anoxic basin: Nitinat Lake depth 20 m clone Ni EU265
EU101198.1 gg_id:237363 Sulfide/oxygen supply ratio predicts outcome competition among sulfur-oxidizing wa EU101
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EU245804.1 gg_id:257757 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-M6-F0 EU245
DQ330597.1 gg_id:159245 Guerrero Negro hypersaline microbial mat clone 09D2Z91 DQ330
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EU245526.1 gg_id:256848 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-M2-F0 EU245
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FJ416124.1 gg_id:357229 sediments sediment station NEC5 Northern Bering Sea clone 115O3 FJ4161
EU542519.1 gg_id:273867 Effects chemical structure concentration on pathways microbial during dechlorinatio EU542
DQ811954.1 gg_id:162191 mangrove soil clone MSB-5A5 DQ811
EU487916.1 gg_id:564815 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU487
EF203188.1 gg_id:204759 phylotypes sediments Lake Kastoria Greece sediment clone Kas139B EF2031
EU048677.1 gg_id:254199 and archaeal surface marine sediment Slope South China Sea clone MD2896-B18 EU048
EU487964.1 gg_id:573020 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU487
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FM165273.1 gg_id:566144 seep chitineous tube vestimentiferan tubeworm Lamellibrachia sp. cold seeps clone FM165
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EU245601.1 gg_id:257385 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-M3-E0 EU245
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FJ901586.1 gg_id:567970 phreatic limestone sinkholes northeastern Mexico biomat 30m deep cenote La Palita FJ9015
AB240362.1 gg_id:143463 microbial structure Phragmites rhizosphere biofilm reed bed reactor laboratory clone AB240
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AJ543759.1 gg_id:95866 anoxic sediment Lake Cadagno clone 49 AJ5437
FJ746261.1 gg_id:567096 Microbial oligotrophic marine sediments: South Pacific Gyre ocean sediment 5306 m FJ7462
AB237732.1 gg_id:139838 microorganisms deep subsurface groundwater sedimentary rock milieu clone HDBW AB237
EU386097.1 gg_id:261462 Biogeographical and microbes sediments on subseafloor sediment South China Sea c EU386
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EF687508.1 gg_id:283688 Biogeochemistry iron- and sulfur-precipitating microbial mats (Nile Mediterranean) s EF6875
GQ249498.1 gg_id:562766 Microbial anaerobic sediments Bay Cadiz (Spain) marine sediment: 0-2 cm layer clor GQ249
DQ329567.1 gg_id:159197 Guerrero Negro hypersaline microbial mat clone 08D2Z50 DQ329
DQ329541.1 gg_id:159505 Guerrero Negro hypersaline microbial mat clone 05D2Z11 DQ329
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DQ329542.1 gg_id:159334 Guerrero Negro hypersaline microbial mat clone 05D2Z93 DQ329
EU407206.1 gg_id:273907 household biogas digester clone GW-22 EU407
EU246196.1 gg_id:246894 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-P5-B01 EU246
FJ484771.1 gg_id:566342 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4847
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FJ712479.1 gg_id:532343 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) k FJ7124
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FJ716279.1 gg_id:551134 Chlorobi dominate shal chemocline Bahamas Sawmill Sink column 10.3 m water dept FJ7162
GU363012.1 gg_id:532990 and archaeal biogeochemical cycling genes within Qiongdongnan basin Sea: pilot stu GU363
AY114321.1 gg_id:63785 anoxic marine sediment clone LD1-PA32 AY114
EU487944.1 gg_id:562517 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU487

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EU373977.1 gg_id:275456 southern Cretan margin (eastern Mediterranean Sea) canyon and slope sediment clc EU373
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EU236367.1 gg_id:299835 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg91E2 EU236
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EU885059.1 gg_id:323370 microbial hindgut chamber Pomacanthus sexstriatus GI tract clone A2F3 EU885
EU491122.1 gg_id:266959 Abundance and microbial life ocean crust seafloor lavas Loi'hi Seamount Pisces Peak EU491
U81755.2 gg_id:1542 anaerobic digestor clone vadinHB65 U8175
AB009200.1 gg_id:14576 rumen clone RFN44 AB009
EU491130.1 gg_id:265874 Abundance and microbial life ocean crust seafloor lavas Loi'hi Seamount Pisces Peak EU491
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EF092196.1 gg_id:274447 Persistent Axinella corrugata Lentisphaerae clone Ax29_B12 EF0921
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EU462436.1 gg_id:290815 Sumatran orangutan feces clone orang1_aai54e09 EU462
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GU180194.1 gg_id:547075 Impact coplanar PCBs on microbial activity anaerobic estuarine sediments sediment GU180
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EF106489.1 gg_id:202958 microbial two endoevaporitic salt crusts evaporitic crust Guerrero Negro Baja clone 2 EF1064
AJ133617.1 gg_id:1688 brine sediment clone KTK 32 AJ1336
EU246019.1 gg_id:246314 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-P1-F02 EU246
FN393496.1 gg_id:552766 prokaryotic sediments two Tunisian multipond sediment solar saltern clone SFD1H1 FN393
GU390826.1 gg_id:526942 Dark hot springs: structure biofilms caves Acquasanta Terme (Italy) sulfidic cave stre GU390
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CU921965.1 gg_id:554874 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU921
AY570581.1 gg_id:108459 -temperature biodegraded Canadian oil reservoir clone PL-11B10 AY570
GU476613.1 gg_id:515032 microbial biogas fermentation enrichment clone BBMC-13 GU476
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DQ354746.1 gg_id:152989 subsurface water clone MS149BH1062003_13 DQ354
GU196193.1 gg_id:591769 Bioaugmentation improved recovery digesters after toxicant exposure lab-scale ana GU196
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FJ748807.1 gg_id:578422 Vertical and sulfate-reducing prokaryotes estuarine Southern China Pearl River Estua FJ7488
AJ519650.1 gg_id:75874 uranium mill tailings soil sample clone Sh765B-TzT-35 AJ5196
AB426199.1 gg_id:274653 -situ identification study benzene-degrading s under methanogenic condition anaer AB426
AB486305.1 gg_id:538794 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSBAR AB486
EF492972.1 gg_id:257902 Iron-manganese Nodule and Its Surrounding Soil soil around iron-manganese nodule EF4929
AF351214.1 gg_id:71271 coal tar-contaminated groundwater clone 12-24 AF351
FJ621558.1 gg_id:561816 Enrichment and detection denitrifying methanotrophic NC10 phylum eutrophic fresh FJ6215
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FJ907182.1 gg_id:573993 Enrichment anoxic methane utilizing denitrification (AnMUD) microorganisms mixed FJ9071
DQ369742.1 gg_id:156874 microbial consortium couples oxidation denitrification anaerobic methane oxidation DQ369
AB179508.1 gg_id:105296 sedimentary rock clone MIZ17 AB179
DQ906849.1 gg_id:250973 Oman subsurface soil clone 10A-11 DQ906
DQ906843.1 gg_id:244437 Oman subsurface soil clone 7C-23 DQ906
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AM991185.1 gg_id:278069 Microbial groundwater and their potential use biomonitoring karst spring water clo AM991
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EU373968.1 gg_id:273635 southern Cretan margin (eastern Mediterranean Sea) canyon and slope sediment cl EU373
EU335184.1 gg_id:242324 Changes microbial metabolic and along hydrogeochemically variable profile saturate EU335
AF351231.1 gg_id:42281 upgradient coal tar waste-contaminated groundwater clone 4-29 sp. AF351
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DQ138963.1 gg_id:138815 rice straw paddy soil clone JG118 DQ138

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AF544226.1 gg_id:58150 acid mine drainage clone ASL9 AF5442
AB254780.1 gg_id:159816 Biogenic Fe-oxyhydroxides nodules (takashikozo) encrusted iron-oxide reducing sedi AB2547
EU372645.1 gg_id:564896 Evaluation Leptospirillum spp. Rio Tinto basin model interest biohydrometallurgy wa EU3726
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FJ228338.1 gg_id:347059 sulfate-reducing enriched during bioremediation experiment sediments (Brandenburg FJ2283
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EF409845.1 gg_id:211316 acid mine drainage clone FKBZ-53-06 EF4098
AF543509.1 gg_id:56204 acid mine drainage clone BA33 AF5435
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AF155154.1 gg_id:1493 Nitrospira cf. moscoviensis str. SBR2016 AF1551
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AY555798.1 gg_id:106808 Bor Khlueng hot spring clone PK287 AY5557
EU287071.1 gg_id:237590 Pacific arctic surface sediment clone B78-107 EU2870
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EU803576.1 gg_id:332256 It's all ranking aquatic Lake Gatun clone 5C231165 EU8035
GQ249372.1 gg_id:583840 Research on biodiversity sludge bioflim Biological Aeration Filter clone B14 GQ249
DQ814725.1 gg_id:173004 zebrafish digestive tract clone aaa35h01 DQ814
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AY921743.1 gg_id:114282 farm soil clone AKYG586 AY9217
EU491464.1 gg_id:276859 Abundance and microbial life ocean crust seafloor lavas Hawai'i South Point X4 clone EU4914
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GU118642.1 gg_id:549595 Threatened Corals Provide Microbial Habitats clone Mfav_D14 GU118
EF492939.1 gg_id:257467 Iron-manganese Nodule and Its Surrounding Soil soil around iron-manganese nodule EF4929
GQ848453.1 gg_id:588843 Microbial sediments vents along East Lau Spreading Centre deep sea hydrothermal GQ848
GQ500804.1 gg_id:586905 Surveys Biofilms and Sediments Mammoth Cave Karst Aquifers Bio-Sep beads conta GQ500
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AY921494.2 gg_id:153641 Changes soils rice field soil clone AP19 AY9214
AY493911.1 gg_id:136740 Fingerprinting aggregates soil clone 1183 AY4939
NR_029287.1 gg_id:554350 Nitrospira moscoviensis str. NSP M-1; DSM 10035 NR_02
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EU652665.1 gg_id:279215 Seasonal variation microbial Yel Sea sediment clone D8S-37 EU652
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EU491363.1 gg_id:274991 Abundance and microbial life ocean crust seafloor lavas Hawai'i South Point X3 clone EU491
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DQ354748.1 gg_id:155453 subsurface water clone MS149BH1062003_38 DQ354
FJ484454.1 gg_id:540847 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4844
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DQ811894.1 gg_id:171789 mangrove soil clone MSB-3B5 DQ811
DQ833491.1 gg_id:167800 detected analyses depth sediment Guanting Reservoir clone 35-52 DQ833
FJ902311.1 gg_id:558620 phreatic limestone sinkholes northeastern Mexico biomat rock outcrop cenote La Pal FJ9023
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FJ497634.1 gg_id:570634 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4976
FJ497567.1 gg_id:539167 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4975
GQ246342.1 gg_id:566822 Temporal and spatial China North Yel Sea sediments clone Cm1-5 GQ246
EU617857.1 gg_id:306314 Seasonal variation microbial Yel Sea sediment clone C13S-3 EU617
FJ769506.1 gg_id:514019 Changes Methanogenic Microbial After Sludge Pre-treatment full-scale municipal ana FJ7695
EU245141.1 gg_id:238328 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-H2-H0 EU245
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EU491069.1 gg_id:263387 Abundance and microbial life ocean crust seafloor lavas Loi'hi Seamount South Rift X EU491
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EU591656.1 gg_id:275968 Microbial Ecology Focused-Pulsed Electric Field Sludge Pretreatment Methanogenic EU591
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DQ067040.1 gg_id:132726 Lake Washington sediment clone pLW-99 DQ067
GU118009.1 gg_id:554824 Threatened Corals Provide Microbial Habitats clone Apal_G06 GU118
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EU635936.1 gg_id:278503 Microbiology geochemistry great boiling and mud springs sediment hot spring; temp EU635
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EU924242.1 gg_id:310156 Microbiology and geochemistry Little Hot Creek hot spring sediment temperature 8 EU9242
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EU050864.1 gg_id:253554 libraries sediment Kings Bay Svalbard Arctic clone SS1_B_08_10 EU0508
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EU386085.1 gg_id:258248 Biogeographical and microbes sediments on subseafloor sediment South China Sea c EU386
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EU800499.1 gg_id:331078 It's all ranking aquatic Delaware Bay NJ clone 2C228620 EU800
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AM991275.1 gg_id:279473 Microbial groundwater and their potential use biomonitoring karst spring water clo AM991
DQ676354.1 gg_id:226121 Archaeal sediment and plankton freshwater pond suboxic freshwater-pond clone M DQ676
AM997371.1 gg_id:538271 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
AF507707.1 gg_id:65679 forest soil clone S042 AF507
AY661975.1 gg_id:137070 uranium-contaminated groundwater clone 300E2-C09 AY661
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AM997579.1 gg_id:552101 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
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DQ837261.1 gg_id:238430 Archaeal and pristine National Park Spain Donana coastal aquifer clone 49S1_2B_70 DQ837
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EF019262.1 gg_id:204198 aspen rhizosphere clone Elev_16S_444 EF0192
AB294344.1 gg_id:211556 Microbial structure deep coal seem groundwater stream clone SWB33 AB294
EU881347.1 gg_id:317758 (maize-sweet cropping field) Karst region maize-sweet potato cropland soil clone KM EU881
EU491266.1 gg_id:274405 Abundance and microbial life ocean crust seafloor lavas Loi'hi Seamount Pisces Peak EU491
EU373974.1 gg_id:272286 southern Cretan margin (eastern Mediterranean Sea) canyon and slope sediment clc EU373
FJ901639.1 gg_id:509494 phreatic limestone sinkholes northeastern Mexico biomat 30m deep cenote La Palita FJ9016
FJ901656.1 gg_id:593533 phreatic limestone sinkholes northeastern Mexico biomat 30m deep cenote La Palita FJ9016
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EU385872.1 gg_id:263088 Stratified microbes sediments Core MD05-2902 subseafloor sediment South China S EU385
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EU385795.1 gg_id:269229 Stratified microbes sediments Core MD05-2896 subseafloor sediment South China S EU385
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AF449263.1 gg_id:61847 Riftia pachyptila's tube clone R76-B102 AF4492
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EF092217.1 gg_id:270621 Persistent Axinella corrugata clone B255_19a EF0922
FJ208822.1 gg_id:333159 Enrichment ANAMMOX Wastewater Treatment Plants Korea and Their Activity sludg FJ2088
FJ208814.1 gg_id:336106 Enrichment ANAMMOX Wastewater Treatment Plants Korea and Their Activity sludg FJ2088
FJ208815.1 gg_id:318107 Enrichment ANAMMOX Wastewater Treatment Plants Korea and Their Activity sludg FJ2088
EF516837.1 gg_id:223894 grassland soil clone FCPS655 EF5168
GQ349982.1 gg_id:579975 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ349
DQ513105.1 gg_id:161200 Microbial life fluids ridge flank crustal fluid clone FS275-24B-03 DQ513
EU373996.1 gg_id:270915 southern Cretan margin (eastern Mediterranean Sea) canyon and slope sediment clc EU373
EU491188.1 gg_id:266276 Abundance and microbial life ocean crust seafloor lavas Loi'hi Seamount Pisces Peak EU491
EU478635.1 gg_id:332383 microdiversity study anammox Candidatus Scalindua phylotype marine oxygen minir EU478
EF645945.1 gg_id:242961 Potential interactions particle- anammox and archaeal partners namibian Namibian u EF6459
GU118095.1 gg_id:584707 Threatened Corals Provide Microbial Habitats clone Apal_E05 GU118
EU050872.1 gg_id:237785 libraries sediment Kings Bay Svalbard Arctic clone SS1_B_04_05 EU050
DQ070801.1 gg_id:147446 Prokaryotic Structures Seafloor Basalts Varying Age and Juan de Fuca Ridge basalt gl DQ070
CU917688.1 gg_id:569653 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU917
AM056027.1 gg_id:137533 anammox wastewater treatment plant AM056
AM490671.1 gg_id:550928 Linking and functional nutrient spiraling mats (USA) microbial mat sulfidic cave sprir AM490
AY266450.1 gg_id:85294 Baltimore Harbor sediment clone B4 AY2664
EU478641.1 gg_id:338211 microdiversity study anammox Candidatus Scalindua phylotype marine oxygen minir EU478
FN396716.1 gg_id:565403 constant flux thermophilic cold seabed Arctic marine surface sediment clone s5_8_II FN3967
EU135126.1 gg_id:257290 tallgrass prarie soil clone FFCH12905 EU1351
FJ444691.1 gg_id:565666 Cotton Rhizosphere cotton rhizosphere clone 3h-166 FJ4446
FJ905739.1 gg_id:535979 s Tonga-Kermadec iron oxide sediments Volcano 1 Tonga Arc clone V1F106b FJ9057
GQ339140.1 gg_id:527968 Comprehensive Investigation on Iron Cycling Freshwater Seep ing Microscopy Cultiv GQ339
EU617888.1 gg_id:353402 Seasonal variation microbial Yel Sea sediment clone D13S-58 EU6178
EU638793.1 gg_id:340317 Ecological Role Firmicutes Identified Thermophilic Microbial Fuel Cells thermophilic n EU6387
GQ347619.1 gg_id:580887 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
AF314425.1 gg_id:39505 aerobic phosphorus-removal ecosystem clone PHOS-HE93 AF3144
EU478694.1 gg_id:312765 microdiversity study anammox Candidatus Scalindua phylotype marine oxygen minir EU478
FJ710640.1 gg_id:513938 Abundance and microbial long-term operated anammox reactors initiated inocula upf FJ7106
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GQ356125.1 gg_id:576581 biomarkers ammonia oxidation activity anaerobic ammonium-oxidation (Anammox) GQ356
EU661858.1 gg_id:292847 and kinetic research ammonia mixed sludge anaerobic ammonium oxidation reactor EU6618
AB057453.1 gg_id:28149 Planctomycete clone KSU-1 AB0574
GQ175284.1 gg_id:578161 nitrogen removal reactors lab scale SBR inorganic medium and glucose influence org GQ175
AY254883.1 gg_id:102606 rotating biological contactor clone EN 8 AY2548
AY257181.1 gg_id:92388 Candidatus Scalindua sorokinii AY2571
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EF602038.1 gg_id:296196 Anammox marine sediments: cell numbers potential anammox activity and pore nutr EF6020

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EU142947.1 gg_id:324638 Enrichment and two marine anammox involved nitrogen gas production anaerobic a EU142
AB300486.1 gg_id:219357 Detection Activity and Anammox Genes Coastal Marine Sediments Japan intertidal s AB300
AB281489.1 gg_id:194829 anaerobic ammonium-oxidizing planctomycete JMK-2 AB281
EU478593.1 gg_id:324123 microdiversity study anammox Candidatus Scalindua phylotype marine oxygen minir EU478
EU039861.1 gg_id:316215 Anaerobic Ammonium Oxidation Anammox Oxygen Minimum Zone Northern upper EU039
EU478626.1 gg_id:345186 microdiversity study anammox Candidatus Scalindua phylotype marine oxygen minir EU478
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EU478595.1 gg_id:335981 microdiversity study anammox Candidatus Scalindua phylotype marine oxygen minir EU478
DQ869384.1 gg_id:173040 evidence broad anaerobic ammonium-oxidizing freshwater and marine sediments B DQ869
EU039841.1 gg_id:338086 Anaerobic Ammonium Oxidation Anammox Oxygen Minimum Zone Northern upper EU039
EU478687.1 gg_id:306841 microdiversity study anammox Candidatus Scalindua phylotype marine oxygen minir EU478
EU039858.1 gg_id:318377 Anaerobic Ammonium Oxidation Anammox Oxygen Minimum Zone Northern upper EU039
EU039856.1 gg_id:305926 Anaerobic Ammonium Oxidation Anammox Oxygen Minimum Zone Northern upper EU039
GQ143786.1 gg_id:381880 Level Continental Shelf Sediment and Phylum-level lineages Phylum Planctomycetes GQ143
EU478661.1 gg_id:328021 microdiversity study anammox Candidatus Scalindua phylotype marine oxygen minir EU478
EU039855.1 gg_id:347502 Anaerobic Ammonium Oxidation Anammox Oxygen Minimum Zone Northern upper EU039
EU478664.1 gg_id:307650 microdiversity study anammox Candidatus Scalindua phylotype marine oxygen minir EU478
DQ534739.1 gg_id:229398 Anaerobic ammonium oxidation contributes significantly nitrogen loss Peruvian oxy DQ534
EU385873.1 gg_id:258876 Stratified microbes sediments Core MD05-2902 subseafloor sediment South China S EU385
EU385800.1 gg_id:262268 Stratified microbes sediments Core MD05-2896 subseafloor sediment South China S EU385
AB177204.1 gg_id:131205 seafloor methane hydrate clone ODP1230B31.20 AB177
EU050874.1 gg_id:239076 libraries sediment Kings Bay Svalbard Arctic clone SS1_B_03_21 EU050
AB474364.1 gg_id:554768 Phycisphaera mikrensis str. NBRC 102668 AB474
FJ516779.1 gg_id:569736 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled biofilm FJ5167
EU236426.1 gg_id:302876 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg92H6 EU236
FJ202723.1 gg_id:356606 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2027
EU491354.1 gg_id:261113 Abundance and microbial life ocean crust seafloor lavas Hawai'i South Point X3 clone EU491
AF445715.1 gg_id:44762 Planctomycetales clone SM2B12 AF445
DQ811901.1 gg_id:173711 mangrove soil clone MSB-4A3 DQ811
EF632950.1 gg_id:252359 Unique microbial contrasting aquatic altitude Andean Altiplano (northern Chile) fresh EF6329
AY344414.1 gg_id:91792 Lake Kauhako surface clone K2-S-7 AY344
FJ205228.1 gg_id:533393 Prokaryotic deep-sea region East Lau Spreading Centre inactive hydrothermal field se FJ2052
EU135058.1 gg_id:239668 tallgrass prarie soil clone FFCH1059 EU135
FJ205225.1 gg_id:568694 Prokaryotic deep-sea region East Lau Spreading Centre inactive hydrothermal field se FJ2052
EU801770.1 gg_id:341049 It's all ranking aquatic Chesapeake Bay MD clone 3C003141 EU801
FJ626886.1 gg_id:510050 Identification Phytodetritus Degrading Microbial Sublittoral Gulf Mexico Sands ocean FJ6268
EU135079.1 gg_id:247437 tallgrass prarie soil clone FFCH3889 EU135
AB430337.1 gg_id:276470 Removal Nitrogen Compounds Coupling Anoxic Ammonium Oxidation and Nitrate Re AB430

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EU236371.1 gg_id:294838 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg9F4 EU236
AM086132.1 gg_id:145659 Abundance and archaea mesotrophic Lake Kinneret lake profundal sediment clone (AM086
FJ612210.1 gg_id:561975 and Biogeochemistry Dongping Lake Ecosystems lake water clone DP7.3.10 FJ6122
DQ396220.1 gg_id:151370 deep-sea octacoral clone ctg_NISA377 DQ396
DQ513014.1 gg_id:160569 Microbial life fluids ridge flank crustal fluid clone FS140-16B-02 DQ513
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EF574227.1 gg_id:231309 site S25 near Coco's Island marine clone S25_571 EF5742
EF632831.1 gg_id:237452 Unique microbial contrasting aquatic altitude Andean Altiplano (northern Chile) sedir EF6328
AY957928.1 gg_id:115016 drinking water biofilm clone B3NR69D13 AY957
EU135121.1 gg_id:248077 tallgrass prarie soil clone FFCH5834 EU135
DQ450805.1 gg_id:200837 Microbial Alpine Tundra Wet Meadow Soil: Chloroflexi Cold Water-Saturated saturat DQ450
EU662552.1 gg_id:553506 Productivity- Chemolithoautotrophically Based Sulfidic Karst Systems floating microb EU662
EF515938.1 gg_id:218719 grassland soil clone FCPP573 EF515
FJ901683.1 gg_id:591480 phreatic limestone sinkholes northeastern Mexico biomat sediment cenote La Palita (FJ9016
GU359065.1 gg_id:531410 rhizosphere soil peanut clone 1S21 GU359
AF407717.1 gg_id:201735 microbial Australia's Great Artesian Basin 52 deg. C brown filament clone B16 AF407
GQ356133.1 gg_id:566719 biomarkers ammonia oxidation activity anaerobic ammonium-oxidation (Anammox) GQ356
EU881297.1 gg_id:304737 (maize-sweet cropping field) Karst region maize-sweet potato cropland soil clone KM EU881
AF312220.1 gg_id:199817 -based BTX-catabolizing microbial African sandy soil South Africa: KwaZulu-Natal Pro AF312
EF580964.1 gg_id:277693 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM1 EF5809
AF121889.1 gg_id:200587 analyses sulfidogenic and phenol-degrading microbial consortia 2-bromophenol-deh AF1218
DQ521520.1 gg_id:202207 Antarctic lake ice cover clone ANTLV2_E09 DQ521
AJ616270.1 gg_id:202670 and seasonal changes planctomycetales biofilms river biofilm clone DEL26 AJ6162
AB240380.1 gg_id:201941 microbial structure Phragmites rhizosphere biofilm reed bed reactor laboratory clon AB240
EF516039.1 gg_id:225108 grassland soil clone FCPN712 EF5160
EF125955.1 gg_id:581157 Biofilm-reactor Based Enrichment Utilizing Oligotrophic Concentrations Acyl-homose EF1259
FJ232661.1 gg_id:308368 Enrichment ANAMMOX Wastewater Treatment Plants Korea and Their Activity sludg FJ2326
AB486233.1 gg_id:540991 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSBARC AB486
EU135254.1 gg_id:255557 tallgrass prarie soil clone FFCH8636 EU135
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EU135178.1 gg_id:237749 tallgrass prarie soil clone FFCH7061 EU135
FJ232680.1 gg_id:333924 Enrichment ANAMMOX Wastewater Treatment Plants Korea and Their Activity sludg FJ2326
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FJ479427.1 gg_id:586523 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4794
EF580956.1 gg_id:279355 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM9 EF5809
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EU135175.1 gg_id:248027 tallgrass prarie soil clone FFCH6996 EU135
EU044117.1 gg_id:239077 luences structure fosters richness subtropical Altamaha Ohopee River Valley inland c EU044

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EU881267.1 gg_id:321885 (long-term grazing and burned annual) Karst region serious disturbance field soil clor EU881
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FJ466217.1 gg_id:563036 and carbon monoxide-oxidizing across successional on recent Hawaiian 1959 Kilauea FJ4662
FJ536899.1 gg_id:590906 Linking performance microbiology biofilters treating dimethyl sulphide presence and FJ5368
FJ203423.1 gg_id:331911 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2034
EU135107.1 gg_id:249175 tallgrass prarie soil clone FFCH2726 EU135
FJ232673.1 gg_id:325828 Enrichment ANAMMOX Wastewater Treatment Plants Korea and Their Activity sludge FJ2326
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FJ652504.1 gg_id:593106 aerobic and anaerobic ammonia-oxidizing sponges marine sponge clone ML-Pla-38 FJ6525
EF019897.1 gg_id:205038 aspen rhizosphere clone Elev_16S_1264 EF0198
EU881193.1 gg_id:326719 Karst region natural restoration land soil clone KNR200711-121 EU881
EU135245.1 gg_id:250454 tallgrass prarie soil clone FFCH17044 EU135
AJ231195.1 gg_id:2884 Isosphaera pallida AJ2311
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DQ814734.1 gg_id:200402 Reciprocal Gut Microbiota Transplants Zebrafish and Mice Germ-free Recipients Hos DQ814
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GQ402671.1 gg_id:552820 Accessing microbial biodiversity and metabolic potential tropical -based approach p GQ402
AB238776.1 gg_id:146076 shal subsurface Sarobetsu Mire peat soil clone HSM-SS-013 AB238
GQ264465.1 gg_id:545566 Structure Simulated Level Waste Site simulated level waste site clone WW3_167 GQ264
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EU289510.1 gg_id:246083 metagenomic approach biosynthesis ansamycin endophytes or symbionts enriched EU289
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EF515969.1 gg_id:220676 grassland soil clone FCPP710 EF5159
AB254788.1 gg_id:160585 Biogenic Fe-oxyhydroxides nodules (takashikozo) encrusted iron-oxide reducing sedi AB254
AJ867920.1 gg_id:202171 Fluctuations Significantly Influence Microbial Present Nival Lakes lake water clone C8 AJ8679
EU861926.1 gg_id:345379 effects chronic fertilization on alpine tundra microbial implications carbon and cyclin EU861
EF019037.1 gg_id:215283 aspen rhizosphere clone Amb_16S_1586 EF0190
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AF465649.1 gg_id:35108 planctomycete clone YNPRH1A AF4656
EF516492.1 gg_id:222936 grassland soil clone FCPS404 EF5164
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AF234700.1 gg_id:35556 sludge clone A25 AF2347
GQ356088.1 gg_id:542965 biomarkers ammonia oxidation activity anaerobic ammonium-oxidation (Anammox) GQ356
EF516829.1 gg_id:219667 grassland soil clone FCPN696 EF5168
EU135135.1 gg_id:235394 tallgrass prarie soil clone FFCH1662 EU135
EU589315.1 gg_id:273690 China Ningxia rice paddy field soil clone 2_H12 EU5893
AF234760.1 gg_id:33376 sludge clone A17 AF2347
FJ232659.1 gg_id:322102 Enrichment ANAMMOX Wastewater Treatment Plants Korea and Their Activity sludg FJ2326
FJ543069.1 gg_id:531452 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone B FJ5430
X86391.1 gg_id:2815 Pirellula sp. str. AGA/M12 X86391
GQ264007.1 gg_id:509308 Structure Simulated Level Waste Site simulated level waste site clone WC3_03 GQ264
M34126.1 gg_id:2826 Pirellula staleyi M3412
AM902592.1 gg_id:245810 Acidobacteria and Planctomycetes radioactive mineral Central Alps Austria subsurfa AM902
AJ616292.1 gg_id:107831 river biofilm clone DEL77 AJ6162
EU135081.1 gg_id:257073 tallgrass prarie soil clone FFCH2281 EU135
AJ252628.1 gg_id:28765 agricultural soil clone SC-I-28 AJ2526
AF513967.1 gg_id:62476 planctomycete clone LA4-B40N AF5139
X62911.1 gg_id:2845 Planctomyces limnophilus X62911
GQ850548.1 gg_id:568227 surface water northern Bering Sea clone u78 GQ850
AJ231186.1 gg_id:2859 Planctomyces sp. str. 248 AJ2311
EF019893.1 gg_id:207109 aspen rhizosphere clone Elev_16S_1259 EF0198
AJ616272.1 gg_id:106693 river biofilm clone DEL30 AJ6162
EU236333.1 gg_id:288195 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg5a2A6 EU2363
AB286452.1 gg_id:205500 Reducing production and domination comamonadaceae reducing oxygen supply was AB2864
EU703485.1 gg_id:385507 lakes on Eastern Tin Plateau freshwater lake altitude 4279 masl clone XZZLH76 EU7034
GQ264049.1 gg_id:532634 Structure Simulated Level Waste Site simulated level waste site clone WC3_146 GQ264
EU135201.1 gg_id:246619 tallgrass prarie soil clone FFCH11725 EU135
EU135291.1 gg_id:243965 tallgrass prarie soil clone FFCH3333 EU135
FJ232636.1 gg_id:318175 Enrichment ANAMMOX Wastewater Treatment Plants Korea and Their Activity sludg FJ2326
GU118118.1 gg_id:577627 Threatened Corals Provide Microbial Habitats clone Apal_L16 GU118
GQ093884.1 gg_id:415267 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ093
DQ256391.1 gg_id:143227 Cloning and microorganism genes alkaline soil clone pGXAR2 DQ256
EF580936.1 gg_id:278198 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM4 EF5809
GQ202636.1 gg_id:576162 RFLP two library insect-resistant trans-genic and its control soil conventional cotton GQ202
FJ232624.1 gg_id:325188 Enrichment ANAMMOX Wastewater Treatment Plants Korea and Their Activity sludg FJ2326
FJ236055.1 gg_id:356355 three Parisian treatment plants approaches drinking water clone Joinville4 FJ2360
AJ576406.1 gg_id:105661 midgut homogenate Pachnoda ephippiata larva clone PeM40 AJ5764
GQ264445.1 gg_id:591299 Structure Simulated Level Waste Site simulated level waste site clone WW3_146 GQ264
AJ231185.1 gg_id:2863 Planctomyces sp. str. 423 AJ2311

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GQ487909.1 gg_id:585391 Microbial metal soil polluted heavy metals clone T1-58 GQ487
DQ228417.1 gg_id:199905 microbial compositions two subglacial possible role microbes chemical weathering p DQ228
EU979049.1 gg_id:308618 Effects intercropping Rhizobium inoculation on microbial biomass and rhizosphere b EU979
AJ252614.1 gg_id:46917 agricultural soil clone SC-I-11 AJ2526
EF157243.1 gg_id:255306 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF1572
GQ488002.1 gg_id:533542 Microbial metal soil polluted heavy metals clone V8-4 GQ488
FJ624888.1 gg_id:509799 Influence lead contamination on boreal pine forest soil clone HF_C_15 FJ6248
GQ994843.1 gg_id:588567 planctomycete bloom eutrophic and simulated blooms algal microcosms lake sedim GQ994
FJ542938.1 gg_id:517269 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone B FJ5429
EU491846.1 gg_id:270163 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU491
GQ487907.1 gg_id:589189 Microbial metal soil polluted heavy metals clone T1-42 GQ487
FJ437980.1 gg_id:545573 Benthic okenone production Fayetteville New York Green Lake surface sediments 16. FJ4379
EU617893.1 gg_id:314251 Seasonal variation microbial Yel Sea sediment clone D13S-21 EU617
DQ815011.1 gg_id:162552 zebrafish digestive tract clone aab56g07 DQ815
FJ542891.1 gg_id:521367 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone A FJ5428
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GU183590.1 gg_id:556104 Structures Reduced and Oxidized Aquifer lower Aqueous Arsenic Titas Bangladesh a GU183
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AJ616260.1 gg_id:101336 river biofilm clone DEL04 AJ6162
BX294907.2 gg_id:84917 anaerobic digester clone CY0ARA056H03 BX2949
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AY328798.1 gg_id:88168 drinking water system simulator clone DSSD101 AY3287
FJ230918.1 gg_id:339412 under Level Antibiotic Selection Pressures river water clone F45 FJ2309
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EF421451.1 gg_id:215170 Rhodopirellula baltica str. OJF13 EF4214
EF589343.1 gg_id:227672 Rhodopirellula baltica str. OJF16 EF5893
EF012748.1 gg_id:192974 Rhodopirellula baltica str. OJF4 EF0127
EF012749.1 gg_id:180732 Rhodopirellula baltica str. OJF5 EF0127
EU236437.1 gg_id:297164 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg63H3 EU2364
AM259894.1 gg_id:212998 Spatial sponge- Mediterranean Tethya aurantia sponge cortex clone TAA-10-09 AM259
EU236342.1 gg_id:295035 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg5a2H2 EU2363
FJ624356.1 gg_id:440418 Rhodopirellula sp. str. SM50 FJ6243
DQ640663.1 gg_id:233756 full-scale EBPR sludge clone Skagenf17 DQ640
FJ624322.1 gg_id:448778 Rhodopirellula sp. str. CS14 FJ6243

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X81942.1 gg_id:2801 Pirellula sp. str. 302 X81942
EU249951.1 gg_id:245163 Polyphasic description Pocillopora seawater adjacent Pacillopora meandrina coral co EU2499
EF157256.1 gg_id:245908 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF1572
FJ624353.1 gg_id:418512 Rhodopirellula sp. str. SM44 FJ6243
DQ986201.1 gg_id:190032 Pirellula sp. OJF3 DQ986
EF012750.1 gg_id:184921 Rhodopirellula baltica str. OJF6 EF0127
X81939.1 gg_id:2767 Pirellula sp. str. 140 X81939
EU812984.1 gg_id:323488 Microbial structure thermophilic reactors biogas reactor fed corn silage clone TUM-M EU8129
DQ499312.1 gg_id:159626 acidic cave snottite biofilm clone CV73 DQ499
GQ246290.1 gg_id:512628 Temporal and spatial China North Yel Sea sediments clone Sd1-9 GQ246
EU246802.1 gg_id:240923 polyphasic description Pocillopora meandrina Palmyra Atoll Calcinus obscurus abdon EU2468
AJ616261.1 gg_id:101508 river biofilm clone DEL05 AJ6162
AY907796.1 gg_id:136912 identification picoplankton contrasting waters Arabian Sea water clone A714001 AY9077
AB294955.1 gg_id:222378 -dependent & analyses hydrothermal system off Taketomi Island Japan microbial ma AB2949
GU061671.1 gg_id:549851 and function microbial mesoscale eddy perturbations South China Sea clone CE2-DC GU061
DQ903990.1 gg_id:235788 Microbes Hawaiian Marine Sponge Suberites zeteki USA: Oahu Hawaii clone PRPR39 DQ903
X81940.1 gg_id:2783 Pirellula sp. str. 516 X81940
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EF574895.1 gg_id:233369 site S25 near Coco's Island marine clone S25_1239 EF5748
FJ623363.1 gg_id:583343 Efficient polyhydroxyalkanoates production waste-activated submitted process seque FJ6233
FJ624354.1 gg_id:395015 Rhodopirellula sp. str. SM48 FJ6243
GU061406.1 gg_id:569946 and function microbial mesoscale eddy perturbations South China Sea clone CE1-5m GU061
EU037330.2 gg_id:269075 structure landfill sediments receiving long-term augmentations wastes chromium co EU0373
X81943.1 gg_id:2800 Pirellula sp. str. 382 X81943
X81941.1 gg_id:2787 Pirellula sp. str. 158 X81941
X81945.1 gg_id:2793 Pirellula sp. str. 139 X81945
FJ624364.1 gg_id:421926 Rhodopirellula sp. str. SM23 FJ6243
GQ302560.1 gg_id:542353 prokaryotic sediment Shawan methods cold spring clone sw-xj213 GQ302
EU703306.1 gg_id:435220 lakes on Eastern Tin Plateau polysaline lake altitude 3204 masl clone XZQH106 EU7033
AJ431347.1 gg_id:40268 Greenland:Ikka Fjord clone AJ4313
AF468237.1 gg_id:80863 Arctic pack ice; northern Fram Strait; 80 31.1 N; 01 deg 59.7 min E clone ARKCH2Br2-9 AF4682
EU491585.1 gg_id:271404 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR4C EU4915
FJ624347.1 gg_id:400480 Rhodopirellula baltica str. SH155 IFAM 1436 FJ6243
FJ205295.1 gg_id:557224 Prokaryotic deep-sea region East Lau Spreading Centre active hydrothermal field sedi FJ2052
EU050866.1 gg_id:237562 libraries sediment Kings Bay Svalbard Arctic clone SS1_B_06_87 EU0508
FJ485080.1 gg_id:560755 microbial autonomous rover world's deepest phreatic sinkhole biomat El Zacaton 114 FJ4850
FJ197610.1 gg_id:338655 deep Pacific Ocean and Eastern Mediterranean Sea sediment clone ALVIN29a_5E_FL FJ1976
GQ143793.1 gg_id:403814 Level Continental Shelf Sediment and Phylum-level lineages Phylum Planctomycetes GQ143
GQ339128.1 gg_id:517627 Comprehensive Investigation on Iron Cycling Freshwater Seep ing Microscopy Cultiv GQ339
AB177200.1 gg_id:137395 seafloor methane hydrate clone ODP1230B30.13 AB1772

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AY921874.1 gg_id:114298 farm soil clone AKYG1711 AY9218
GU444061.1 gg_id:523807 Effects amendments on fertility and microbial structure cucumber Cucumis sativus L GU444
DQ206406.1 gg_id:139615 microbial mediates arsenate reduction soda lake water clone ML--10 DQ206
AM997881.1 gg_id:576223 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
EF516155.1 gg_id:227320 grassland soil clone FCPT723 EF5161
AB530247.1 gg_id:544394 Monitoring Suez and Tokyo bays based on analyses marine sediment clone Tokyo..B AB530
AF507705.1 gg_id:65683 forest soil clone S165 AF5077
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FJ746253.1 gg_id:510156 Microbial oligotrophic marine sediments: South Pacific Gyre ocean sediment 5306 m FJ7462
EU048670.1 gg_id:257221 and archaeal surface marine sediment Slope South China Sea clone MD2896-B10 EU048
FJ873309.1 gg_id:562253 Vertical and archaea methane-rich cold seep located base sediments Okhotsk Sea clo FJ8733
EF688380.1 gg_id:237185 soil clone F6-89 EF6883
EU245970.1 gg_id:250462 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-M8-H0 EU2459
GU322886.1 gg_id:548781 Metaproteomics -temperature anaerobic granular sludge cultivated temperatures c GU322
GQ246434.1 gg_id:594302 Temporal and spatial China North Yel Sea sediments clone Ld1-39 GQ246
AY493903.1 gg_id:136324 Fingerprinting aggregates soil clone 1430 AY4939
FJ264747.1 gg_id:311487 Manganese and Iron Dependent Anaerobic Oxidation Methane methane seep sedim FJ2647
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GQ246448.1 gg_id:576923 Temporal and spatial China North Yel Sea sediments clone Ld1-53 GQ246
AF507704.1 gg_id:55354 forest soil clone C167 AF5077
AY963300.1 gg_id:156060 structures and correlated ecological factors soils two evergreen broad-leaved forests AY9633
EU386080.1 gg_id:262812 Biogeographical and microbes sediments on subseafloor sediment South China Sea c EU386
FJ479214.1 gg_id:543366 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4792
U81656.2 gg_id:2794 anaerobic digestor clone vadinBA30 U8165
EF019225.1 gg_id:210216 aspen rhizosphere clone Elev_16S_394 EF0192
FJ712480.1 gg_id:518323 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) K FJ7124
FJ638603.1 gg_id:576295 Temperature dependent succession sulfate reducing muddy southwestern Taiwan hc FJ6386
EU622299.1 gg_id:296920 syntrophic partnerships deep-sea vents direct cell capture and metagenomics Eel Riv EU622
EF019958.1 gg_id:215591 aspen rhizosphere clone Elev_16S_1337 EF0199
EU245568.1 gg_id:248646 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-M3-B0 EU2455
EU135296.1 gg_id:244874 tallgrass prarie soil clone FFCH4400 EU135
EU135033.1 gg_id:235601 tallgrass prarie soil clone FFCH5199 EU135
EF516309.1 gg_id:223655 grassland soil clone FCPT556 EF5163
EU245782.1 gg_id:251795 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-M6-C0 EU2457
FJ264744.1 gg_id:327743 Manganese and Iron Dependent Anaerobic Oxidation Methane methane seep sedim FJ2647
EU385736.1 gg_id:271883 Stratified microbes sediments Core MD05-2896 subseafloor sediment South China S EU3857
AF423302.1 gg_id:35667 soil clone 96-2 AF4233
EU245905.1 gg_id:258094 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-M8-A0 EU2459
DQ329791.1 gg_id:159475 Guerrero Negro hypersaline microbial mat clone 05D2Z14 DQ329
EF612394.1 gg_id:255351 historic semiarid lead-zinc mine tailings site soil clone OS-C18 EF6123

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EU385723.1 gg_id:262361 Stratified microbes sediments Core MD05-2896 subseafloor sediment South China S EU385
CU919054.1 gg_id:548383 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU919
EU786130.1 gg_id:556144 cotton rhizosphere soil clone 3H-211 EU786
EF018222.1 gg_id:203959 aspen rhizosphere clone Amb_16S_572 EF0182
EU385921.1 gg_id:264090 Stratified microbes sediments Core MD05-2902 subseafloor sediment South China S EU385
EU385781.1 gg_id:268863 Stratified microbes sediments Core MD05-2896 subseafloor sediment South China S EU385
FJ485338.1 gg_id:571346 microbial autonomous rover world's deepest phreatic sinkhole wall biomat El Zacator FJ4853
EF540419.1 gg_id:220924 oil-shale semi-coke solid waste clone MK11 EF5404
EF032768.1 gg_id:181593 cyanobacterial mat in lava cave clone HAVOmat68 EF0327
EU245501.1 gg_id:245089 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-M2-D0 EU245
AB177155.1 gg_id:136191 seafloor methane hydrate clone ODP1230B16.23 AB177
GQ844339.1 gg_id:543377 Microbial full-scale produced water treatment plant biofilm anoxic tank clone BB-B5 GQ844
GQ472311.1 gg_id:573896 Microbial Dongping Lake lake sediment clone 2B3-12 GQ472
FJ748808.1 gg_id:585249 Vertical and sulfate-reducing prokaryotes estuarine Southern China Pearl River Estua FJ7488
GQ143768.1 gg_id:454105 Level Continental Shelf Sediment and Phylum-level lineages Phylum Planctomycetes GQ143
EU386008.1 gg_id:273669 Biogeographical and microbes sediments on subseafloor sediment South China Sea c EU386
DQ811900.1 gg_id:163946 mangrove soil clone MSB-5C12 DQ811
FJ545451.1 gg_id:508778 Diveristy Seawater and Sediment Northern North Yel Sea sediment clone S1-19 FJ5454
EU245488.1 gg_id:235811 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-M2-B0 EU245
EF208700.1 gg_id:200413 Vertical and horizontal structure microbial sandy carbonate sediment clone CI75cm.2 EF2087
EF516637.1 gg_id:225058 grassland soil clone FCPS673 EF5166
AJ292685.1 gg_id:44216 PCB-polluted soil clone WD283 AJ2926
FJ444683.1 gg_id:588682 Cotton Rhizosphere cotton rhizosphere clone 2y-55 FJ4446
GU056117.1 gg_id:554003 Biologic Techniques Applied Microbial Prospecting China: Ban 876 Gas Oil Field Bein GU056
FN563400.1 gg_id:556620 Impact changing HRT OLR on prokaryotes involved biomethanizitaion and fuzzy logic FN563
EF018678.1 gg_id:211992 aspen rhizosphere clone Amb_16S_968 EF0186
EF515994.1 gg_id:219383 grassland soil clone FCPT539 EF5159
EU181513.1 gg_id:237389 Prokaryotic enhanced organic matter remineralization continental margin marine se EU181
EF019563.1 gg_id:214259 aspen rhizosphere clone Elev_16S_1077 EF0195
EU491712.1 gg_id:272697 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU491
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FJ712491.1 gg_id:510851 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) k FJ7124
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FJ478514.1 gg_id:544225 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4785
FJ232658.1 gg_id:353016 Enrichment ANAMMOX Wastewater Treatment Plants Korea and Their Activity sludg FJ2326
EU491293.1 gg_id:271990 Abundance and microbial life ocean crust seafloor lavas Loi'hi Seamount Pisces Peak EU491
EU135104.1 gg_id:243304 tallgrass prarie soil clone FFCH17847 EU135
EU374014.1 gg_id:271620 southern Cretan margin (eastern Mediterranean Sea) canyon and slope sediment clc EU374
AY592336.1 gg_id:104089 deep-sea mud volcano clone Amsterdam-1B-32; BC20-1B-32 AY5923
FJ203528.1 gg_id:337295 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2035

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FJ873317.1 gg_id:512846 Vertical and archaea methane-rich cold seep located base sediments Okhotsk Sea clo FJ8733
GQ356148.1 gg_id:571807 biomarkers ammonia oxidation activity anaerobic ammonium-oxidation (Anammox) GQ356
FJ516887.1 gg_id:535958 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled upper s FJ5168
EU843071.1 gg_id:348461 metagenomics random sampling pyrosequencing rumen fluid fiber adherent microbi EU843
FJ652501.1 gg_id:582419 aerobic and anaerobic ammonia-oxidizing sponges marine sponge clone ML-Pla-33 FJ6525
AY114326.1 gg_id:71342 anoxic marine sediment clone LD1-PA40 AY1143
EU491864.1 gg_id:259028 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU4918
FJ203614.1 gg_id:306565 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2036
FJ203304.1 gg_id:323269 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2033
FJ232644.1 gg_id:337297 Enrichment ANAMMOX Wastewater Treatment Plants Korea and Their Activity sludge FJ2326
DQ396219.1 gg_id:152458 deep-sea octacoral clone ctg_NISA190 DQ396
EF573103.1 gg_id:229876 Coco's Island site 23 marine clone S23_1202 EF5731
GQ487912.1 gg_id:578134 Microbial metal soil polluted heavy metals clone T1-103 GQ487
FJ205299.1 gg_id:548823 Prokaryotic deep-sea region East Lau Spreading Centre active hydrothermal field sedi FJ2052
FJ203110.1 gg_id:337594 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2031
EU734992.1 gg_id:341990 assemblages sediment station DBSE Northern Bering Sea clone 086B35 EU734
FJ203078.1 gg_id:355875 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2030
GQ356114.1 gg_id:512540 biomarkers ammonia oxidation activity anaerobic ammonium-oxidation (Anammox) GQ356
AB486712.1 gg_id:589011 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSCOR AB486
EF123574.1 gg_id:249219 Microbial disease microbial mats on Siderastrea siderea three regions wider Caribbea EF1235
EF125439.1 gg_id:179213 mangrove soil clone MSB-2E1 EF1254
GQ340148.1 gg_id:586939 Prokaryotic Marathonas Reservoir Greece water column clone B07-89-BAC GQ340
AM774187.1 gg_id:278832 lineages Planctomycetes densely colonize alkaline gut soil-feeding termites (Cubiter AM774
AB192132.1 gg_id:202625 termite gut homogenate clone M1NP1-64 AB192
AM086149.1 gg_id:149208 Abundance and archaea mesotrophic Lake Kinneret lake profundal sediment clone AM086
GU369884.1 gg_id:518971 and mats vents Arc white filamentous microbial mat photic shal hydrothermal vent GU369
AF234735.1 gg_id:50882 sludge clone A8 AF2347
EF020031.1 gg_id:216224 aspen rhizosphere clone Elev_16S_1426 EF0200
CU926663.1 gg_id:535989 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU926
EF125452.1 gg_id:190111 mangrove soil clone MSB-2F5 EF1254
GU118783.1 gg_id:558104 Threatened Corals Provide Microbial Habitats clone Mfra_K07 GU118
GQ340267.1 gg_id:579129 Prokaryotic Marathonas Reservoir Greece water column clone A08-139-BAC GQ340
AJ306767.1 gg_id:66932 DCP-dechlorinating consortium clone SHA-87 AJ3067
FJ203428.1 gg_id:327231 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2034
EF516721.1 gg_id:218584 grassland soil clone FCPU585 EF5167

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DQ329768.1 gg_id:201435 Guerrero Negro hypersaline microbial mat clone 02D2Z12 DQ329
EU386107.1 gg_id:265536 Biogeographical and microbes sediments on subseafloor sediment South China Sea c EU386
DQ070794.1 gg_id:147997 Prokaryotic Structures Seafloor Basalts Varying Age and Juan de Fuca Ridge basalt gl DQ070
FJ203440.1 gg_id:321503 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2034
AB486437.1 gg_id:552968 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSBARC AB486
GU302676.1 gg_id:533316 inhabiting epithelial tissue ruminal digesta -DGGE and sequencing rumen contents c GU302
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EU135213.1 gg_id:244630 tallgrass prarie soil clone FFCH13721 EU135
GQ214098.1 gg_id:562881 China loess clone P818 GQ214
EF540354.1 gg_id:225493 oil-shale semi-coke solid waste clone 24B_MJK EF5403
AY704401.1 gg_id:112006 oceanic crust clone FS142-21B-02 AY704
FJ232639.1 gg_id:332829 Enrichment ANAMMOX Wastewater Treatment Plants Korea and Their Activity sludge FJ2326
GU118848.1 gg_id:532921 Threatened Corals Provide Microbial Habitats clone Mfra_N03 GU118
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DQ404806.1 gg_id:155344 contaminated sediment clone 661261 DQ404
EF205581.1 gg_id:214356 among geographically regions central Tibet geothermal spring mat clone TP5 EF2055
GU118276.1 gg_id:548121 Threatened Corals Provide Microbial Habitats clone Dstr_C07 GU118
FJ232651.1 gg_id:351648 Enrichment ANAMMOX Wastewater Treatment Plants Korea and Their Activity sludge FJ2326
DQ404633.1 gg_id:152343 contaminated sediment clone 656078 DQ404
EF125406.1 gg_id:178940 soil naked tidal flat sediment near mangrove clone MSB-1E4 EF1254
FJ203554.1 gg_id:342344 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2035
EF208715.1 gg_id:200049 Vertical and horizontal structure microbial sandy carbonate sediment clone CI75cm.2 EF2087
DQ289931.1 gg_id:170286 Microbial Carbon and Nitrogen Cycling Sediments South Atlantic Bight Permeable Sh DQ289
EF208689.1 gg_id:200911 Vertical and horizontal structure microbial sandy carbonate sediment clone CI5cm.FC EF2086
AB240225.1 gg_id:142901 microbial structure Phragmites rhizosphere biofilm reed bed reactor laboratory clone AB240
AB270056.1 gg_id:170512 Influence temperature and humidity on heifers rumen contents Holstein heifer (Bos AB270
AJ290182.1 gg_id:200064 and seasonal changes planctomycetales biofilms extracted river biofilm s clone DSP2! AJ2901
EU381635.1 gg_id:264273 Rumen eubacterial implicates Bacteroidetes major particulate phylum rumen liquid p EU381
EF419414.1 gg_id:204219 Substantial alterations cutaneous biota psoriatic lesions human skin clone 6P2-75 EF4194
FJ516944.1 gg_id:509690 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled upper s FJ5169
FJ542842.1 gg_id:525666 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone A FJ5428
X81948.1 gg_id:2823 Pirellula str. 670 X81948
GQ246406.1 gg_id:536385 Temporal and spatial China North Yel Sea sediments clone Ld1-11 GQ246
DQ289943.1 gg_id:168811 Microbial Carbon and Nitrogen Cycling Sediments South Atlantic Bight Permeable Sh DQ289
EU488313.1 gg_id:555142 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU488
FJ232657.1 gg_id:342256 Enrichment ANAMMOX Wastewater Treatment Plants Korea and Their Activity sludge FJ2326
AY225652.1 gg_id:96259 Mid-Atlantic Ridge hydrothermal sediment clone AT-s2-38 AY2256
FJ437939.1 gg_id:544528 Benthic okenone production Fayetteville New York Green Lake surface sediments 16. FJ4379
DQ070834.1 gg_id:150611 Prokaryotic Structures Seafloor Basalts Varying Age East Pacific Rise and Ridge basal DQ070

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FJ542996.1 gg_id:514874 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone B FJ5429
EU734986.1 gg_id:353483 assemblages sediment station DBSE Northern Bering Sea clone 080B24 EU7349
EU491549.1 gg_id:269000 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR4C EU4915
FJ172792.1 gg_id:344848 Qinghai Yak Ruminal and Fungi China rumen clone RB-2C12 FJ1727
FJ203088.1 gg_id:312705 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2030
EF494313.1 gg_id:224266 Nunnock River granitic landscape clone NR.4.006 EF4943
DQ984569.1 gg_id:245832 Mt. Nan-Jen soil clone IFD_124 DQ984
AB355065.1 gg_id:252268 Monitoring and archaeal two heavy polluted areas lake Egypt based on sediment Ma AB3550
EF515912.1 gg_id:217349 grassland soil clone FCPO648 EF5159
AB487699.1 gg_id:588740 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB4876
FJ208839.1 gg_id:351437 Enrichment ANAMMOX Wastewater Treatment Plants Korea and Their Activity sludge FJ2088
GQ402727.1 gg_id:545813 Accessing microbial biodiversity and metabolic potential tropical -based approach pe GQ402
FJ905699.1 gg_id:513469 s Tonga-Kermadec iron oxide sediments Volcano 1 Tonga Arc clone V1F13b FJ9056
FJ652511.1 gg_id:586795 aerobic and anaerobic ammonia-oxidizing sponges marine sponge clone ML-Pla-24 FJ6525
FJ208828.1 gg_id:315617 Enrichment ANAMMOX Wastewater Treatment Plants Korea and Their Activity sludge FJ2088
EU621897.1 gg_id:274014 and kinetic research ammonia mixed anaerobic ammonium oxidation reactor seedec EU6218
CU920107.1 gg_id:537364 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU920
FJ374260.1 gg_id:359089 Inter- and intraspecific humivorous scarab beetle larvae (Pachnoda spp.) larval hindgu FJ3742
AB234544.1 gg_id:145313 termite gut homogenate clone MgMjD-104 AB2345
AB234542.1 gg_id:149255 termite gut homogenate clone MgMjR-038 AB2345
EF032767.1 gg_id:178484 cyanobacterial mat in lava cave clone HAVOmat22 EF0327
AB252879.1 gg_id:171614 Identification Shibayama lagoon Ishikawa Japan iron-oxidation biofilm clone 018 AB2528
EU881130.1 gg_id:329424 Karst region primary forest soil clone KPF200711-199 EU8811
EU135304.1 gg_id:244428 tallgrass prarie soil clone FFCH773 EU1353
GU119349.1 gg_id:579236 Threatened Corals Provide Microbial Habitats reef water clone Reef_L13 GU119
GQ348014.1 gg_id:555348 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
EF032766.1 gg_id:189073 cyanobacterial mat in lava cave clone HAVOmat101 EF0327
EU617890.1 gg_id:309328 Seasonal variation microbial Yel Sea sediment clone D13S-65 EU6178
EU135137.1 gg_id:242995 tallgrass prarie soil clone FFCH15608 EU1351
EU478637.1 gg_id:324136 microdiversity study anammox Candidatus Scalindua phylotype marine oxygen minir EU4786
AY947993.1 gg_id:143867 Synchrony and seasonality two temperate rivers USA: Massachusetts Parker River cl AY9479
AJ247193.1 gg_id:25580 Asticcacaulis biprosthecium str. DSM 4723(T) AJ2471
AB016610.1 gg_id:5563 Asticcacaulis excentricus str. ATCC15261 AB0166
AF296688.1 gg_id:23972 Brevundimonas alba str. H12C5 AF2966
FJ950569.1 gg_id:436320 Brevundimonas diminuta str. c82 FJ9505
EU536433.1 gg_id:281870 profile human microbiota antecubital fossa (inner elbow) skin clone nbt213g08 EU5364
FJ658350.1 gg_id:568324 Metalworking Fluids and Aerosols Metallurgy Plant metalworking aerosol clone Winte FJ6583
DQ335215.1 gg_id:145289 Brevundimonas terrae str. KSL-145 DQ335
EU465068.1 gg_id:294281 African elephant feces clone AFYEL_aaj67e11 EU4650
FJ950592.1 gg_id:424138 Brevundimonas diminuta str. c94 FJ9505

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EU465065.1 gg_id:301063 African elephant feces clone AFYEL_aaj67e08 EU4650
EU434576.1 gg_id:262199 Brevundimonas diminuta str. b108 EU4345
EU464720.1 gg_id:297700 Hartmann's mountain zebra feces clone AFZEB_aaj69f09 EU4647
FJ658310.1 gg_id:535144 Metalworking Fluids and Aerosols Metallurgy Plant metalworking aerosol clone Winte FJ6583
FJ658395.1 gg_id:540941 Metalworking Fluids and Aerosols Metallurgy Plant metalworking aerosol clone Winte FJ6583
EU434570.1 gg_id:265057 Brevundimonas diminuta str. b77 EU4345
EU464980.1 gg_id:290745 African elephant feces clone AFYEL_aaj66b08 EU4649
DQ824709.1 gg_id:183605 human fecal clone RL185_aaj72b02 DQ824
FJ657857.1 gg_id:537717 Metalworking Fluids and Aerosols Metallurgy Plant metalworking fluid clone Winter_ FJ6578
EU535556.1 gg_id:284254 profile human microbiota antecubital fossa (inner elbow) skin clone nbt27b12 EU5355
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FJ658136.1 gg_id:543663 Metalworking Fluids and Aerosols Metallurgy Plant metalworking aerosol clone Winte FJ6581
EU434575.1 gg_id:258759 Brevundimonas diminuta str. b124 EU4345
AB021415.1 gg_id:5595 Brevundimonas diminuta str. IAM 12691T AB0214
EU465159.1 gg_id:296763 African elephant feces clone AFYEL_aaj68h07 EU4651
FN433035.1 gg_id:559314 Brevundimonas terrae CCM24B FN4330
EU434509.1 gg_id:268353 Brevundimonas diminuta str. b215 EU4345
EU464952.1 gg_id:291623 African elephant feces clone AFYEL_aaj65f11 EU4649
AJ244706.1 gg_id:109577 Brevundimonas mediterranea str. V4.BO.18 LMG 22801 = MM_2367 AJ2447
AJ244707.1 gg_id:109696 Brevundimonas mediterranea str. V4.BE.56 LMG 22802 = MM_3001 AJ2447
AM402981.1 gg_id:162149 Brevundimonas sp. str. JO26 AM402
AY971369.1 gg_id:115361 Brevundimonas kwangchunensis str. KSL-110 AY9713
EU834791.1 gg_id:585677 Microbiology Aerobic Phosphorus Removal Process aerobic phosphorus-removing se EU8347
EF363713.1 gg_id:204302 Brevundimonas lenta str. DS-18 EF3637
GU321351.1 gg_id:547657 Brevundimonas sp. str. NMC11 GU321
AM412000.1 gg_id:198378 Brevundimonas nasdae Z66zhy AM412
AY328801.1 gg_id:90161 drinking water OR9 AY3288
EF157195.1 gg_id:241110 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF1571
AY169433.1 gg_id:82124 Brevundimonas vesicularis AY1694
AJ227782.1 gg_id:5579 Brevundimonas bacteroides str. CB7 AJ2277
AJ227797.1 gg_id:5583 Brevundimonas sp. str. FWC40 AJ2277
AJ227781.1 gg_id:5611 Brevundimonas vesicularis str. LMG 11141 AJ2277
AY293404.1 gg_id:87601 travertine terrace Yellowstone Hot Springs clone SM2A11 AY2934
EF546433.1 gg_id:219465 Brevundimonas nasdae str. P1 EF5464
EU182863.1 gg_id:247068 Brachybacterium sp. str. MH35 EU1828
DQ825663.1 gg_id:162728 Brevundimonas sp. str. 5420S-28 DQ825
AY328842.1 gg_id:89703 drinking water MB11 AY3288
AJ227786.1 gg_id:53033 Brevundimonas intermedia str. ATCC 15262(T) AJ2277
EU636022.1 gg_id:278329 Antarctic str. N97 EU6360
EU182890.1 gg_id:250309 Brevundimonas sp. str. MH62 EU1828

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EU143355.1 gg_id:238339 Brevundimonas sp. str. J22 EU143
EU580513.1 gg_id:543062 Benthic diatoms Constance: functional interactions regarding growth and EPS-secret EU580
AJ007808.1 gg_id:5626 Brevundimonas variabilis str. ACM 5107 (T) AJ0078
AB426562.1 gg_id:270554 Brevundimonas sp. str. HINF004 AB4265
DQ372984.1 gg_id:152214 Brevundimonas sp. str. EMB102 DQ372
AY328835.1 gg_id:89630 drinking water MA26 AY3288
EU379308.1 gg_id:274963 Brevundimonas intermedia str. 6R-J-7d EU3793
AJ007802.1 gg_id:5568 Brevundimonas intermedia str. ACM 2608 (T) AJ0078
AJ227784.1 gg_id:5617 Brevundimonas subvibrioides str. CB81 AJ2277
AJ495803.1 gg_id:63466 Brevundimonas sp. p22P AJ4958
AJ227783.1 gg_id:43186 Brevundimonas variabilis str. ATCC 15255 (T) AJ2277
EU434579.1 gg_id:266190 Brevundimonas aurantiaca str. e12 EU4345
AM183353.1 gg_id:167639 Brevundimonas sp. 1228-1II AM183
AB447548.1 gg_id:316703 Brevundimonas sp. str. SUWAF011 AB4475
DQ981459.1 gg_id:203778 Brevundimonas sp. SB4 DQ981
DQ442520.1 gg_id:191932 Streptomyces longisporoflavus str. NRRL ISP-5165T DQ442
FN397631.1 gg_id:452397 Brevundimonas sp. str. CM 272 FN3976
DQ413152.1 gg_id:152156 Brevundimonas sp. str. EMB 68 DQ413
AJ007803.1 gg_id:5518 Caulobacter fusiformis str. ACM 5108 (T) AJ0078
AB008533.1 gg_id:5557 Caulobacter fusiformis str. ATCC15257 AB0085
AJ007805.1 gg_id:5560 Caulobacter henricii str. ACM 5105 (T) AJ0078
AE005673.1 gg_id:209341 Caulobacter crescentus str. CB15 AE0056
AM921622.1 gg_id:248395 Caulobacter henricii PhyCEm-80 AM921
EF515958.1 gg_id:221595 grassland soil clone FCPT664 EF5159
DQ124680.1 gg_id:154386 Caulobacter tundrae str. Z-0024 DQ124
AF523036.1 gg_id:89365 natural mineral water clone D-1 AF5230
EU730905.1 gg_id:352383 Caulobacter vibrioides str. 156 EU7309
DQ532299.1 gg_id:164042 Geographical Variation Microbial Burden and Spacecraft Assembly Kennedy Space C DQ532
AY512823.1 gg_id:106146 Caulobacter vibrioides str. S13 AY5128
AB271055.1 gg_id:165771 Caulobacter ginsengisoli str. Gsoil 317 AB2710
AM940947.1 gg_id:278226 Caulobacter sp. ECN-2008 AM940
DQ404656.1 gg_id:151101 contaminated sediment clone 654987 DQ404
GQ396801.1 gg_id:538121 Functional unvegetated perhumid recently-deglaciated soils do not correlate shifts s GQ396
FN556568.1 gg_id:518192 Brevundimonas sp. str. JW23.4a FN5565
EU434505.1 gg_id:263790 Brevundimonas bullata str. b350 EU4345
DQ501309.1 gg_id:199553 Seasonal dynamics free-living and particle- four lakes northeastern Germany freshw DQ501
AY234477.1 gg_id:112129 soil isolate Ellin5060 AY2344
FJ197848.1 gg_id:313604 Brevundimonas sp. 39(2008) str. 39 FJ1978
AJ717391.1 gg_id:111099 Phenylobacterium falsum str. AC49 AJ7173
AJ582031.1 gg_id:93768 soil near uranium mill tailings clone JG36-GS-63 AJ5820
AY921824.1 gg_id:114202 farm soil clone AKYG696 AY9218

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AJ292592.1 gg_id:49162 PCB-polluted soil clone WD225 AJ2925
DQ202243.1 gg_id:140164 Enrichment s Type II Methane-Oxidizing Inhabiting Northern Peatlands peatland clo DQ202
GQ402724.1 gg_id:551851 Accessing microbial biodiversity and metabolic potential tropical -based approach p GQ402
AB473899.1 gg_id:554330 sequences endolithic and archaea Weissenstein (white rock) / Albula pass few millim AB4738
D49422.1 gg_id:5594 Brevundimonas diminuta str. IFO 3140 D49422
DQ984595.1 gg_id:240075 Mt. Nan-Jen litterfall clone IYF98 DQ984
GQ402558.1 gg_id:592079 Accessing microbial biodiversity and metabolic potential tropical -based approach p GQ402
DQ177484.1 gg_id:139814 Brevundimonas sp. str. Tibet-IK73 DQ177
EF516954.1 gg_id:226283 grassland soil clone FCPT726 EF5169
AJ292595.1 gg_id:30983 PCB-polluted soil clone WD233 AJ2925
AF289151.1 gg_id:70005 freshwater clone PRD01a003B AF2891
EU730914.1 gg_id:335822 Brevundimonas diminuta str. 207 EU7309
EU445215.1 gg_id:565413 Isolation Weight Forest Topsoil Metagenomic tropical forest topsoil clone R36 EU4452
GU118509.1 gg_id:573233 Threatened Corals Provide Microbial Habitats clone Gven_K20 GU118
DQ532251.1 gg_id:168212 Geographical Variation Microbial Burden and Spacecraft Assembly Johnson Space C DQ532
GU169058.1 gg_id:570254 Biofilms formed on substances: response f conditions and carbon concentrations sy GU169
FJ979845.1 gg_id:402409 Caulobacter sp. str. tsz12 FJ9798
AM697237.1 gg_id:227434 sequencing libraries indoor dust clone BF0001C078 AM697
AY922007.1 gg_id:113237 farm soil clone AKYG412 AY9220
GU118288.1 gg_id:568445 Threatened Corals Provide Microbial Habitats clone Dstr_G22 GU118
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EU202832.1 gg_id:317304 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU2028
EU434454.1 gg_id:264269 Brevundimonas diminuta str. a229 EU4344
DQ177483.1 gg_id:140010 Brevundimonas sp. str. Tibet-IK53 DQ177
GU118656.1 gg_id:575957 Threatened Corals Provide Microbial Habitats clone Mfav_J15 GU118
DQ532201.1 gg_id:171288 Geographical Variation Microbial Burden and Spacecraft Assembly Johnson Space C DQ532
EU881329.1 gg_id:324088 (maize-sweet cropping field) Karst region maize-sweet potato cropland soil clone KM EU8813
EF520395.1 gg_id:235185 structure lakes: what controls ? acid-impacted lake clone ADK-BTe02-51 EF5203
EU784736.1 gg_id:336351 Brevundimonas sp. str. E8 EU7847
AY569007.1 gg_id:103285 Nitrobacteria hamadaniensis str. 104 AY5690
AB167225.1 gg_id:107935 Brevundimonas diminuta str. c66 AB1672
EF520409.1 gg_id:231965 structure lakes: what controls ? acid-impacted lake clone ADK-BTh02-54 EF5204
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GU118170.1 gg_id:560087 Threatened Corals Provide Microbial Habitats clone Dstr_P10 GU118
AB473919.1 gg_id:573162 sequences endolithic and archaea Weissenstein (white rock) / Albula pass few millim AB4739
EU665093.1 gg_id:547599 Soil are Changed Agricultural Practices agricultural soil clone Plot22-C12 EU6650
AB374373.1 gg_id:272478 sequences Weissenstein (white rock) / few millimeters be surface rock called 'Weiss AB3743
AB167229.1 gg_id:107937 Brevundimonas diminuta str. c79 AB1672
AY734246.1 gg_id:133283 spring water of alpine karst aquifer clone SDKAS1_2 AY7342
DQ167245.1 gg_id:138986 Kopriimonas byunsanensis str. KOPRI 13522 DQ167

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AB373113.1 gg_id:277271 Thalassospira sp. str. N5EA13-1A AB3731
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AF358664.1 gg_id:35557 Thalassospira lucentensis AF3586
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AB006768.1 gg_id:4594 Terasakiella pusilla str. IFO13613 AB0067
AY922191.1 gg_id:112927 whalefall clone 131616 AY9221
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AJ633974.1 gg_id:114828 host's accessory nidamental gland clone 6ANG531 AJ6339
EF629835.1 gg_id:258350 Changes marine Mycale laxissima on transfer aqua sponge tissue EF6298
AY922227.1 gg_id:113991 whalefall clone 131732 AY9222
GQ169074.1 gg_id:592716 Thalassospira lucentensis str. TVGB18 GQ169
AY162078.1 gg_id:135758 Cultivating Sargasso Sea clone GMD37B4 AY1620
EU880514.1 gg_id:352101 Thalassospira lucentensis str. P44 EU8805
FJ952806.1 gg_id:399348 Thalassospira sp. str. 2ta2 FJ9528
AF143826.1 gg_id:25696 hydrocarbon-degrading consortium clone 4-Org1-58 AF1438
GU118853.1 gg_id:541338 Threatened Corals Provide Microbial Habitats clone Mfra_C21 GU118
EU268103.1 gg_id:254391 summer Dalian and Liaodong bay determined sea water clone L0603 EU2681
AY682384.1 gg_id:100730 Parvulomonas gwangyangensis str. GW14-5 AY6823
DQ917818.1 gg_id:221237 Microbial profiles bleached and wild-type Muricea elongata Kordiimonas clone BME DQ917
GU119022.1 gg_id:554110 Threatened Corals Provide Microbial Habitats clone Past_K19 GU119
EU287361.1 gg_id:252140 Pacific arctic surface sediment clone S26-61 EU2873
AB537162.1 gg_id:517870 Preparation single species magnetotactic multiple-displacement sediment water we AB5371
X81185.1 gg_id:11274 magnetic coccus MP17 str. TB24 X81185
X61607.1 gg_id:11262 magnetic coccus MP17 X61607
AAAN03000028.1 gg_id:130031 Magnetococcus sp. str. MC-1 AAAN0
DQ785314.1 gg_id:166095 magnetotactic sediments Jiaozhou Bay sediment clone QH-1 DQ785
EF371486.1 gg_id:214607 magnetotactic China Jiaozhou Bay clone M-40 EF3714
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AJ863157.1 gg_id:111324 marine microcosm clone CF24 AJ8631
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FM242322.1 gg_id:318593 sequences sediment clone 63 T9d-oil FM242
EF530680.1 gg_id:226147 Extremely acidic pendulous biofilms Frasassi cave system Italy microbial biofilm clone EF5306
AY193164.1 gg_id:80157 marine tidal mat clone BTM11 AY1931
GU370118.1 gg_id:530853 Rhodovulum sulfidophilum str. P721(3) GU370
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EU442518.1 gg_id:275257 Aurantimonas altamirensis str. NML 070722 LSPQ no. ID089387 EU4425
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EF373540.1 gg_id:206133 Aurantimonas kwangyangensis str. CW5 EF3735
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EF489025.1 gg_id:221997 Bacillus sp. str. HB01085 EF4890
FM872541.1 gg_id:368575 occupant source house floor dust clone FA01H06 FM872
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GQ115503.1 gg_id:421082 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw686f02 GQ115
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AB246801.2 gg_id:147120 Bartonella elizabethae str. R-5 AB2468
AB440632.1 gg_id:338719 Bartonella sp. str. Fuji 18-1 AB4406
AB242293.1 gg_id:141840 Bartonella sp. str. Fuji 12-1 AB2422

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AB426635.1 gg_id:259103 Bartonella grahamii str. Cg4263alb AB426
AB426630.1 gg_id:276614 Bartonella grahamii str. Hokkaido 48-1 AB426
AB529507.1 gg_id:594215 Bartonella grahamii str. cg147up AB529
AB529498.1 gg_id:578474 Bartonella grahamii str. Korea 4-1 AB529
AY513504.1 gg_id:103693 Bartonella henselae str. 882_ANT5 AY513
U87831.1 gg_id:5492 aortic heart valve patient with endocarditis clone v9 U8783
Z11684.1 gg_id:5473 Bartonella henselae str. BA-TF Z11684
EU111755.1 gg_id:273803 Bartonella queenslandensis str. AUST/NH5 EU111
EU111749.1 gg_id:270242 Bartonella rattiaustraliensis str. AUST/NH4 EU111
EU111754.1 gg_id:263406 Bartonella queenslandensis str. AUST/NH12 EU111
NC_005955.1 gg_id:129029 Bartonella quintana str. Toulouse NC_00
AJ278187.1 gg_id:25998 Bartonella schoenbuchensis str. R1 AJ2781
AB519061.1 gg_id:550137 Bartonella washoensis str. DR1-1 AB519
AB519065.1 gg_id:548080 Bartonella washoensis str. SR25-1 AB519
AB519062.1 gg_id:548956 Bartonella washoensis str. DR3-1 AB519
EF682085.1 gg_id:331242 'Candidatus rudakovii' detected voles Western Siberia Candidatus Bartonella rudakov EF682
AF293389.1 gg_id:49049 Bartonella capreoli str. IBS 193 AF293
FJ177634.1 gg_id:353598 Bartonella sp. str. HMD-3 FJ1776
EU294521.1 gg_id:244723 Bartonella volans str. FSq-1 EU294
Z70001.1 gg_id:5501 Bartonella sp str. R-phy2 Z70001
EF616480.1 gg_id:249397 Bartonella silvicola str. Cul-9 BAA-1453 EF6164
AB292597.1 gg_id:211030 Bartonella washoensis str. Sb944nv AB292
Z70005.1 gg_id:5475 Bartonella sp str. R-phy1 Z70005
Z70008.1 gg_id:5483 Bartonella sp str. C5rat Z70008
EU111759.1 gg_id:268319 Bartonella coopersplainensis str. AUST/NH20 EU111
AF204274.1 gg_id:46747 Bartonella birtlesii str. 325 AF2042
X89110.1 gg_id:5486 Wolbachia melophagi str. MO6 X89110
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DQ645425.1 gg_id:161188 Bartonella henselae str. M7SHD DQ645
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EU111752.1 gg_id:260102 Bartonella rattiaustraliensis str. AUST/NH14 EU111
AY515119.1 gg_id:100507 Bartonella phoceensis str. 16120 AY515
NR_029368.1 gg_id:541643 Bartonella doshiae str. R18; NCTC 12862 NR_02
AY254309.1 gg_id:104973 Bartonella chomelii AY2543
L01260.1 gg_id:5502 Bartonella elizabethae L01260
DQ538394.1 gg_id:227943 Bartonella australis str. Aust/NH1 DQ538
AY515118.1 gg_id:100352 Bartonella rattimassiliensis str. 16115 AY515
AJ002139.1 gg_id:5462 Bartonella alsatica str. 382 AJ0021
NR_029366.1 gg_id:571234 Bartonella grahamii str. V2; NCTC 12860 NR_02

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AB119197.1 gg_id:106044 Beijerinckia indica str. ATCC 9036 AB119
AB119199.1 gg_id:106358 Beijerinckia indica str. ATCC 19361 AB119
AJ563932.1 gg_id:100451 Beijerinckia mobilis str. DSM 2326 AJ5639
AB119210.1 gg_id:106197 Beijerinckia sp. str. TB16 AB119
AB119203.1 gg_id:101664 Beijerinckia sp. str. TB9 AB119
EF584507.1 gg_id:232942 Chelatococcus daeguensis str. K106 EF5845
AM162441.1 gg_id:155035 and situ identification acidic sphagnum peat bog clone B56 AM162
AJ563926.1 gg_id:104947 Methylocella palustris str. Ch3 AJ5639
CP001280.1 gg_id:527956 Methylocella silvestris str. BL2 CP001
AJ555244.1 gg_id:83530 Methylocella tundrae str. T4 AJ5552
DQ450165.1 gg_id:155167 (Uruguay) Atlantic coastal lagoon clone R7C54 DQ450
EU131038.1 gg_id:252176 alkaline China coal mine soil clone SIP CM24 EU131
GQ402592.1 gg_id:584864 Accessing microbial biodiversity and metabolic potential tropical -based approach p GQ402
FJ719048.1 gg_id:530044 Protozoan grazers alter metabolically active aquifer aquifier sediments clone p03_D0 FJ7190
EF064161.1 gg_id:225511 Microbial sediment bottom AMD pool acid mine drainage clone Sed3 EF0641
X97076.1 gg_id:4762 peat bog clone TM9 X97076
EU133271.1 gg_id:246401 tallgrass prarie soil clone FFCH2375 EU133
AF407384.1 gg_id:27469 MCB-contaminated groundwater-treating reactor clone RB7C5 AF4073
FM253682.1 gg_id:348953 Rock gold mine unusually microbial biofilm clone H09-2 FM253
AF200697.1 gg_id:32333 13C- extracted 13C-methanol exposed soil clone UP3 proteobacterium AF2006
AB119198.1 gg_id:106192 Beijerinckia derxii str. ATCC 49361 AB119
FM253655.1 gg_id:349812 Rock gold mine unusually microbial biofilm clone E02-2 FM253
FN433469.1 gg_id:562500 Methylocapsa sp. str. KYG FN4334
CU926994.1 gg_id:551719 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU926
DQ837242.1 gg_id:252587 Archaeal and pristine National Park Spain Donana coastal aquifer clone 49S1_2B_23 DQ837
GU205295.1 gg_id:524604 unusual Microbial Venezuelan Tepui (Orthoquartzite) Cave: Dominance Chloroflexal GU205
AY913393.1 gg_id:112985 forest soil clone DUNssu187 (-7A) (OTU#034) AY9133
DQ451456.1 gg_id:156677 forest soil clone FAC17 DQ451
U87759.1 gg_id:4967 Afipia broomeae str. F186 U87759
AY513503.1 gg_id:103555 Afipia felis str. 2002033830 AY5135
U87764.1 gg_id:4964 Afipia genosp. 2 str. G8965 U87764
EU499727.1 gg_id:276997 Field-based laboratory stable isotope probing surveys identities both aerobic and an EU499
GQ032540.1 gg_id:375131 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ032
DQ222224.1 gg_id:140520 Bradyrhizobium sp. str. CCBAU 71283012 DQ222
AY568510.1 gg_id:102699 Afipia massiliensis str. NW-12 AY5685
AY029562.1 gg_id:58038 Afipia massiliensis str. 34633; CIP 107022; CCUG 45153 AY0295
AY547290.1 gg_id:99065 Bradyrhizobium sp. str. CCBAU 7128301 AY5472
AF288304.1 gg_id:41002 Afipia birgiae str. 34632 AF2883
AB099659.1 gg_id:77914 Oligotropha carboxidovorans str. S23 AB0996
FJ193001.1 gg_id:304744 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1930
FJ165544.1 gg_id:559211 Microbially-enhanced Dissolution Karst Carbonates microbial biofilm Sonterra #8 well FJ1655

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U87761.1 gg_id:4971 Afipia broomeae str. G5958 U8776
AY568503.2 gg_id:106979 Afipia broomeae str. C-12 AY5685
EF203409.1 gg_id:261125 Bradyrhizobium sp. str. URBM1 EF2034
EF520400.1 gg_id:231834 structure lakes: what controls ? acid-impacted lake clone ADK-BTh02-40 EF5204
EU308573.1 gg_id:243951 Deinococcus sp. str. 4A4 EU3085
DQ838529.1 gg_id:161809 Methylobacterium sp. str. WSM3686 DQ838
AJ871433.1 gg_id:111252 Chelatococcus asaccharovorans CP141b AJ8714
DQ298340.1 gg_id:151224 studies on and nifH Lasiurus sindicus Sewan grass Thar Desert Rajasthan India rhizos DQ298
FM160634.1 gg_id:292268 Microvirga sp. PCA19 PCA 19 FM160
AB098515.2 gg_id:78887 Balneimonas flocculans str. TFB AB0985
GQ103792.1 gg_id:446977 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ103
AY776209.1 gg_id:108377 Methylobacterium sp. str. AC72a AY7762
AF338176.1 gg_id:68816 Blastobacter denitrificans str. IFAM 1005; LMG 8443 AF3381
X66025.1 gg_id:5058 Blastobacter denitrificans str. LMG 8443 X66025
AF288306.1 gg_id:24179 Bosea vestrisii str. 34635 AF2883
GU377117.1 gg_id:526769 Afipia sp. str. sptzw29 GU377
EU325943.1 gg_id:255748 Bosea sp. str. fei-4-9 EU3259
DQ377760.1 gg_id:153195 Bosea sp. PD 19 DQ377
AJ968692.1 gg_id:165499 Bosea sp. str. STM 358 AJ9686
AJ250796.1 gg_id:4822 Bosea thiooxidans str. DSM 9653 AJ2507
DQ104985.1 gg_id:136035 Bosea thiooxidans str. JS 11 DQ104
EU730912.1 gg_id:324973 Bosea thiooxidans str. 187 EU7309
AJ250797.1 gg_id:4823 Bosea thiooxidans str. RpPI5-VS AJ2507
DQ303323.1 gg_id:149709 Bosea sp. TM19_7 DQ303
EF114681.1 gg_id:194437 Bosea sp. str. SJH07 EF1146
AF288307.1 gg_id:27298 Bosea massiliensis str. 34649 AF2883
EU130951.1 gg_id:243491 Afipia sp. BAC312 EU1309
AF288309.1 gg_id:47211 Bosea massiliensis str. 63287 AF2883
EU584530.1 gg_id:263418 Afipia sp. str. Everest-gws-91 EU5845
NR_028787.1 gg_id:583653 Bosea minatitlanensis str. AMX51; DSM 13099 NR_028
EU561082.1 gg_id:273777 Bradyrhizobium sp. str. CTFI54 EU5610
FJ390904.1 gg_id:589546 Bradyrhizobium canariense str. SEMIA 928 FJ3909
GQ899183.1 gg_id:569090 Bradyrhizobium canariense str. L-2 GQ899
EU561083.1 gg_id:273925 Bradyrhizobium sp. str. CTFI410 EU5610
EU420082.1 gg_id:569322 Bradyrhizobium canariense str. Cy-1 EU4200
AY649429.1 gg_id:99495 Bradyrhizobium elkanii str. BR2001 AY6494
EU170546.1 gg_id:251540 Bradyrhizobium elkanii str. CCBAA 05065 EU1705
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AY904753.1 gg_id:111860 Bradyrhizobium elkanii str. SEMIA 6148 AY9047

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AY904762.1 gg_id:111745 Bradyrhizobium elkanii str. SEMIA 6160 AY9047
AF417554.1 gg_id:21176 Bradyrhizobium elkanii str. DASA37005 AF4175
AF208516.1 gg_id:55735 Bradyrhizobium elkanii str. USDA 46 AF2085
FJ193395.1 gg_id:328559 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1933
DQ786801.1 gg_id:169640 Bradyrhizobium elkanii str. HaE4 DQ786
FJ193017.1 gg_id:353693 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1930
AF417553.1 gg_id:49609 Bradyrhizobium elkanii str. DASA38001 AF4175
AY904736.1 gg_id:112163 Bradyrhizobium elkanii str. SEMIA 696 AY9047
AY904748.1 gg_id:112153 Bradyrhizobium elkanii str. SEMIA 6100 AY9047
FJ193380.1 gg_id:324669 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1933
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AY039016.1 gg_id:64297 Bradyrhizobium sp. ORS3260 str. ORS 3260 AY0390
AF417555.1 gg_id:21431 Bradyrhizobium elkanii str. DASA35011 AF4175
AY904744.1 gg_id:111950 Bradyrhizobium elkanii str. SEMIA 6028 AY9047
EU170551.1 gg_id:242313 Bradyrhizobium elkanii str. CCBAU 23109 EU1705
AY649433.1 gg_id:103679 Bradyrhizobium elkanii str. BR3279 AY6494
AY904789.1 gg_id:112254 Bradyrhizobium elkanii str. SEMIA 6440 AY9047
FJ192573.1 gg_id:321198 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1925
FJ540914.1 gg_id:586787 Bradyrhizobium yuanmingense str. DTB8 FJ5409
FJ540920.1 gg_id:578473 Bradyrhizobium yuanmingense str. DTC5 FJ5409
AF417549.1 gg_id:20737 Bradyrhizobium japonicum str. DASA24015 AF4175
AY904774.1 gg_id:111573 Bradyrhizobium japonicum str. SEMIA 6319 AY9047
AF193818.1 gg_id:49866 Bradyrhizobium yuanmingense str. B071 AF1938
FJ540960.1 gg_id:585297 Bradyrhizobium yuanmingense str. TSC9 FJ5409
AF417548.1 gg_id:45495 Bradyrhizobium japonicum str. DASA03183 AF4175
EF070133.1 gg_id:196710 Bradyrhizobium yuanmingense str. CCBAU 43022 EF0701
EF061098.1 gg_id:183640 Bradyrhizobium yuanmingense str. CCBAU 45079 EF0610
AY649439.1 gg_id:104594 Bradyrhizobium japonicum str. BR3267 AY6494
AY162036.1 gg_id:135634 Cultivating soil clone PI_GH1.1.B1 AY1620
AF417543.1 gg_id:71773 Bradyrhizobium japonicum str. DASA12006 AF4175
DQ303328.1 gg_id:146763 Dynamic during maturation Tuber magnatum Pico ascoma Bradyrhizobium clone TM DQ303
DQ133341.1 gg_id:137967 Bradyrhizobium liaoningense str. GXD1 DQ133
DQ786798.1 gg_id:168591 Bradyrhizobium liaoningense str. DdE3 DQ786
AB240363.1 gg_id:144998 microbial structure Phragmites rhizosphere biofilm reed bed reactor laboratory clone AB2403
AF239843.2 gg_id:32699 Bradyrhizobium japonicum str. PRY40 AF2398
EF221623.1 gg_id:207870 Bradyrhizobium sp. str. JR008 EF2216

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AY904778.1 gg_id:112228 Bradyrhizobium elkanii str. SEMIA 6387 AY9047
AY904757.1 gg_id:112271 Bradyrhizobium japonicum str. SEMIA 6155 AY9047
DQ303324.1 gg_id:147536 Dynamic during maturation Tuber magnatum Pico ascoma Bradyrhizobium clone TM DQ303
EU145989.1 gg_id:236262 Bradyrhizobium liaoningense str. CCBAU 83684 EU1459
EU170561.1 gg_id:245983 Bradyrhizobium elkanii str. CCBAU 41236 EU1705
EU399713.1 gg_id:258733 Bradyrhizobium japonicum str. CCBAU 43170 EU3997
AF417544.1 gg_id:32002 Bradyrhizobium japonicum str. DASA12007 AF4175
GQ181045.1 gg_id:563588 Bradyrhizobium elkanii str. LN5 GQ181
DQ303376.1 gg_id:145414 Dynamic during maturation Tuber magnatum Pico ascoma Bradyrhizobium clone TM DQ303
AF417542.1 gg_id:23765 Bradyrhizobium japonicum str. DASA02010 AF4175
EU561080.1 gg_id:266287 Bradyrhizobium sp. str. CTFI47 EU5610
FJ025097.1 gg_id:348667 Bradyrhizobium japonicum str. SEMIA 6153 FJ0250
EF394156.1 gg_id:210105 Bradyrhizobium japonicum str. CCBAU 65068 EF3941
FJ390914.1 gg_id:591406 Bradyrhizobium japonicum str. SEMIA 512 FJ3909
AB195987.1 gg_id:132401 Bradyrhizobium elkanii str. MF834 AB1959
X70403.1 gg_id:4945 Bradyrhizobium sp str. LMG 9566 X70403
AB179515.1 gg_id:100954 sedimentary rock clone MIZ24 AB1795
AF363138.1 gg_id:24680 Bradyrhizobium japonicum str. MSDJ G94 AF3631
EF394146.1 gg_id:214964 Bradyrhizobium japonicum str. CCBAU 53135 EF3941
FJ418696.1 gg_id:364095 Bradyrhizobium japonicum str. CCBAU 33175 FJ4186
AY904746.1 gg_id:111411 Bradyrhizobium elkanii str. SEMIA 6069 AY9047
DQ517956.1 gg_id:159686 Bradyrhizobium japonicum str. HHL-01 DQ517
GQ181034.1 gg_id:510301 Bradyrhizobium elkanii str. LNW14 GQ181
EU133476.1 gg_id:237433 tallgrass prarie soil clone FFCH17074 EU1334
EF070138.1 gg_id:195443 Bradyrhizobium liaoningense str. CCBAU 33158 EF0701
DQ303346.1 gg_id:145143 Dynamic during maturation Tuber magnatum Pico ascoma Bradyrhizobium clone TM DQ303
AY904756.1 gg_id:112304 Bradyrhizobium japonicum str. SEMIA 6152 AY9047
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AF239844.2 gg_id:51967 Bradyrhizobium japonicum str. PRY42 AF2398
AY568513.2 gg_id:103141 Bradyrhizobium elkanii str. NW-6 AY5685
DQ303379.1 gg_id:146676 Dynamic during maturation Tuber magnatum Pico ascoma Bradyrhizobium clone TM DQ303
EU335263.1 gg_id:257149 Changes microbial metabolic and along hydrogeochemically variable profile unsatura EU3352
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DQ303364.1 gg_id:148890 Dynamic during maturation Tuber magnatum Pico ascoma Bradyrhizobium clone TM DQ303
EF512281.1 gg_id:223867 Bradyrhizobium japonicum str. CCBAU 45002 EF5122
AF363135.1 gg_id:18027 Bradyrhizobium japonicum str. MSDJ 5697 AF3631
EF394151.1 gg_id:211052 Bradyrhizobium japonicum str. CCBAU 65619 EF3941
DQ786799.1 gg_id:170123 Bradyrhizobium liaoningense str. BqE4 DQ786

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DQ303374.1 gg_id:146501 Dynamic during maturation Tuber magnatum Pico ascoma Bradyrhizobium clone TM DQ303
EU074195.1 gg_id:250401 Bradyrhizobium liaoningense str. CCBAU 03282 EU074
DQ303326.1 gg_id:149079 Dynamic during maturation Tuber magnatum Pico ascoma Bradyrhizobium clone TM DQ303
AF417541.1 gg_id:54256 Bradyrhizobium japonicum str. DASA02007 AF4175
AF363133.1 gg_id:37117 Bradyrhizobium japonicum str. MSDJ 5718 AF363
AY599677.1 gg_id:104113 fungal ascocarp clone Cl-19-TB8-II AY5996
AF417545.1 gg_id:49277 Bradyrhizobium japonicum str. THA7 AF4175
EU333380.1 gg_id:268064 Bradyrhizobium japonicum RLA09 EU3333
EU083497.1 gg_id:240912 Evidence Carboxylation During Degradation Hexadecane Denitrifying Consortium he EU083
DQ303335.1 gg_id:148648 Dynamic during maturation Tuber magnatum Pico ascoma Bradyrhizobium clone TM DQ303
AY904773.1 gg_id:111438 Bradyrhizobium elkanii str. SEMIA 6208 AY9047
EF018750.1 gg_id:211106 aspen rhizosphere clone Amb_16S_1209 EF0187
DQ517958.1 gg_id:160432 Bradyrhizobium sp. str. HMS-O1 DQ517
EF019158.1 gg_id:204406 aspen rhizosphere clone Amb_16S_1805 EF0191
U69637.1 gg_id:4921 Bradyrhizobium lupini str. KM50 90 (USDA 3514) U69637
NC_010725.1 gg_id:289367 Methylobacterium populi str. BJ001 NC_01
AF069958.1 gg_id:111772 Nitrobacter alkalicus str. AN4 AF0699
AF069957.1 gg_id:73775 Nitrobacter alkalicus str. AN2 AF0699
NC_007964.1 gg_id:156705 Nitrobacter hamburgensis str. X14 NC_00
AM292301.1 gg_id:254106 Nitrobacter vulgaris str. BB3 AM292
AY055796.1 gg_id:17737 Nitrobacter winogradskyi str. ATCC 14123 AY0557
AM286382.1 gg_id:220346 Nitrobacter vulgaris str. K55 AM286
AY055798.1 gg_id:56503 Nitrobacter sp. str. KB212 AY0557
EU528247.1 gg_id:284778 Seasonal Dynamics Mudflat Mouth Major Kentucky Lake Reservoir Tributary sedime EU528
L11661.1 gg_id:4840 Nitrobacter winogradskyi str. W L11661
AM286390.1 gg_id:223530 Nitrobacter vulgaris str. 329 AM286
L11662.1 gg_id:4842 Nitrobacter sp str. LL L11662
AF529115.1 gg_id:62406 trichloroethene-contaminated site clone FTLM116 proteobacterium AF5291
AB241411.1 gg_id:223626 Rhodopseudomonas lichen O5 AB2414
AB196356.1 gg_id:109667 Rhodopseudomonas palustris str. LW-RE AB1963
AF487428.1 gg_id:68786 Rhodopseudomonas palustris str. Wai1S1 AF4874
U09829.1 gg_id:4878 marine sediment clone JAP850 Rochelle U09829
X87279.1 gg_id:4854 Rhodopseudomonas palustris str. DSM 126 X87279
EU018457.1 gg_id:252635 Rhodobacter sphaeroides PSB07-19 EU0184
CP000463.1 gg_id:183860 Rhodopseudomonas palustris str. BisA53 CP0004
AB167545.1 gg_id:100949 Rhodopseudomonas palustris str. KP0014 FERMP-19432 AB1675
AJ968691.1 gg_id:166553 Rhodopseudomonas sp. 1416ri str. ORS 1416 AJ9686
CP000301.1 gg_id:153177 Rhodopseudomonas palustris str. BisB18 CP0003
EF515308.1 gg_id:224099 Electricigen Enrichment MFC upf microbial fuel cell anode clone 22f08 EF5153
AB167544.1 gg_id:100793 Rhodopseudomonas palustris str. KP0003 FERMP-19430 AB1675

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AF509907.1 gg_id:111553 Bradyrhizobium sp. str. CMI23 AF5099
AB079634.1 gg_id:62340 Rhodopseudomonas palustris VA2-2 AB079
L11664.1 gg_id:4866 Rhodopseudomonas palustris L11664
DQ126338.1 gg_id:138381 Rhodopseudomonas sp. str. PSB-O DQ126
EU531568.1 gg_id:514717 Rhodopseudomonas palustris str. J EU531
EU018458.1 gg_id:248384 Rhodopseudomonas palustris PSB07-21 EU018
GQ036328.1 gg_id:384524 Topographical and Temporal Human Skin Microbiome skin palm clone nbu362h08c1 GQ036
DQ984578.1 gg_id:245601 Mt. Nan-Jen litterfall clone IYF09 DQ984
DQ497243.1 gg_id:157613 Chelatococcus sp DQ497
GQ421849.1 gg_id:550491 Microvirga sp. str. 5420S-16 GQ421
FJ025099.1 gg_id:308491 Bradyrhizobium japonicum str. SEMIA 6396 FJ0250
AY904750.1 gg_id:112390 Bradyrhizobium japonicum str. SEMIA 6144 AY904
AJ009451.1 gg_id:4819 anaerobic TCB-transforming consortium clone SJA-9 AJ0094
EF121353.1 gg_id:216650 microbiota uninfected and Rickettsia-infected colonized cat fleas Ctenocephalides fel EF1213
EU861833.1 gg_id:309753 effects chronic fertilization on alpine tundra microbial implications carbon and nitrog EU861
FJ478995.1 gg_id:579612 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4789
FJ025100.1 gg_id:309477 Bradyrhizobium japonicum str. SEMIA 6154 FJ0251
GU455366.1 gg_id:523224 Understanding short-chain fatty acids accumulation enhanced alkaline digestion: kir GU455
AM412118.1 gg_id:247879 Chelatococcus sp. str. P-117 AM412
EU256461.1 gg_id:256534 Bradyrhizobium sp. str. CCBAU 85044 EU256
AM990457.1 gg_id:279499 Chelatococcus sp. str. I-21 AM990
DQ303342.1 gg_id:150032 Dynamic during maturation Tuber magnatum Pico ascoma Bradyrhizobium clone TM DQ303
EF394148.1 gg_id:205693 Bradyrhizobium japonicum str. CCBAU 43277 EF3941
X87272.1 gg_id:4980 Bradyrhizobium japonicum str. DSM 30131 T X87272
AF530465.1 gg_id:61023 Bradyrhizobium japonicum JZ1 AF5304
DQ129309.1 gg_id:138424 microarray level air two cities Texas urban aerosol clone AKIW763 DQ129
EU276558.1 gg_id:362293 -DGGE and Approaches Analyze Effect Agricultural Practices on Soil agricultural soil c EU276
AF417550.1 gg_id:48037 Bradyrhizobium japonicum str. DASA01059 AF4175
EF018822.1 gg_id:204655 aspen rhizosphere clone Amb_16S_1300 EF0188
FJ716001.1 gg_id:579890 Diet- gut microbiota Lumbricus rubellus fingerprinting technique and cloning intestina FJ7160
AY345540.1 gg_id:91124 Lake Wai'ele'ele water clone LWSR-28 AY3455
GQ129955.1 gg_id:434250 Genetic inventory 'JPL-SAF' during MSL mission spacecraft assembly clean room floc GQ129
AF417547.2 gg_id:33185 Bradyrhizobium japonicum str. DASA38025 AF4175
GQ389009.1 gg_id:587997 quality deterioration drinking system during serious red water outbreak clone J57 GQ389
DQ432334.1 gg_id:156941 prokaryotic alkaline hypersaline lakes Wadi Natrun Egypt Lake Hamra water clone W DQ432
EF394142.1 gg_id:204550 Bradyrhizobium japonicum str. CCBAU 41267 EF3941
AY649440.1 gg_id:99283 Bradyrhizobium japonicum str. BR3288 AY6494
AB187010.1 gg_id:101203 Suiyo seamount hydrothermal vent water 0.2micro-m filterable fraction clone SSM-M AB187
AB195991.1 gg_id:130287 Bradyrhizobium japonicum str. MH615 AB1959
AJ867909.1 gg_id:143411 Fluctuations Significantly Influence Microbial Present Nival Lakes lake water clone B6 AJ8679

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EU878006.2 gg_id:576097 Salinarimonas rosea str. YIM YD3 EU878
GQ103050.1 gg_id:397383 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ103
FJ479132.1 gg_id:512302 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4791
AM292299.1 gg_id:242856 Nitrobacter hamburgensis str. Y AM292
AB195986.1 gg_id:128373 Bradyrhizobium canariense str. MH547 AB195
EU029133.1 gg_id:244029 Deinococcus sp. str. 5A4 EU029
AM411919.1 gg_id:253138 Bradyrhizobium sp. str. P-44 AM411
AB300992.1 gg_id:223242 Bradyrhizobium iriomotense str. EK05 (= NBRC 102520) AB300
EF394144.1 gg_id:214300 Bradyrhizobium japonicum str. CCBAU 53152 EF3941
AY963315.1 gg_id:154614 structures and correlated ecological factors soils two evergreen broad-leaved forests AY9633
AB426223.1 gg_id:263163 -situ identification study benzene-degrading s under methanogenic condition anaerc AB426
AJ785294.1 gg_id:211869 Rhizobiales genosp. AL ARRI 185 AJ7852
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DQ351924.1 gg_id:173105 manganese oxidation soil clone JH-GY18 DQ351
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AY372184.1 gg_id:94343 Bradyrhizobium betae str. PL7HG1 AY372
EF516928.1 gg_id:222831 grassland soil clone FCPN762 EF5169
AF208509.1 gg_id:56515 Bradyrhizobium japonicum str. USDA 129 AF2085
AY904764.1 gg_id:112297 Bradyrhizobium japonicum str. SEMIA 6163 AY904
FJ382130.1 gg_id:582075 Potentially pathogenic and air stem cell transplant summer hospital shower water BM FJ3821
AF417551.1 gg_id:45827 Bradyrhizobium japonicum str. DASA37026 AF417
AY904758.1 gg_id:112060 Bradyrhizobium japonicum str. SEMIA 6156 AY904
FJ528273.1 gg_id:573764 Bradyrhizobium sp. str. Mf253 FJ5282
AF234889.2 gg_id:77279 Bradyrhizobium japonicum str. SEMIA 5080 AF2348
AY904765.1 gg_id:112064 Bradyrhizobium japonicum str. SEMIA 6164 AY904
FJ382683.1 gg_id:510366 Potentially pathogenic and air stem cell transplant summer hospital shower water BM FJ3826
AY904743.1 gg_id:111821 Bradyrhizobium japonicum str. SEMIA 6002 AY904
AF502220.1 gg_id:55851 EBPR sludge lab scale clone HP1B78 AF5022
EF032655.1 gg_id:198655 structure riverine biofilms river biofilm clone SepB-6 EF0326
FJ380996.1 gg_id:549917 Bradyrhizobium sp. str. BQN4T-01d FJ3809
AY178079.1 gg_id:100322 Bradyrhizobium sp. Cmey1 Cmey 1 AY178
AY904732.1 gg_id:111515 Bradyrhizobium japonicum str. SEMIA 656 AY904
AF409021.1 gg_id:98929 Bradyrhizobium sp. str. Ellin179 AF409
D13430.1 gg_id:4937 Bradyrhizobium japonicum str. USDA 110 D13430
AB220164.1 gg_id:153145 Bradyrhizobium sp. QC1 AB220
AF508112.1 gg_id:66234 Bosea thiooxidans TJ1 AF508
GU377100.1 gg_id:514335 Balneimonas sp. str. sptzw09 GU377
AY169423.1 gg_id:81409 Bradyrhizobium japonicum AY1694
D78366.1 gg_id:5055 Agromonas oligotrophica str. JCM1494 D78366
GQ389186.1 gg_id:582505 quality deterioration drinking system during serious red water outbreak clone D59 GQ389

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AF417546.1 gg_id:68929 Bradyrhizobium japonicum str. DASA03066 AF417
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AY904786.1 gg_id:111454 Bradyrhizobium japonicum str. SEMIA 6420 AY904
AM990460.1 gg_id:277676 Chelatococcus sp. str. TKE5A AM990
L37584.1 gg_id:5189 Brucella canis L37584
X13695.1 gg_id:5204 Brucella melitensis biovar Abortus str. 11-19 X13695
EU165529.1 gg_id:255878 Ochrobactrum sp. str. MI9.3 EU165
U70978.1 gg_id:5195 Ochrobactrum anthropi str. LMG 3301 U70978
AF526513.2 gg_id:57487 Ochrobactrum intermedium ADV14 AF5265
EU352765.1 gg_id:265442 Ochrobactrum anthropi str. NK 2.T-3 EU352
AB120120.1 gg_id:94870 Ochrobactrum anthropi AB120
AB091758.1 gg_id:59527 Ochrobactrum tritici AS1 AB091
AM490617.1 gg_id:247109 Ochrobactrum anthropi str. DSM 7216 AM490
FJ984528.1 gg_id:451696 Ochrobactrum haemophilum str. NBSL23 FJ9845
AF229848.1 gg_id:31117 Ochrobactrum sp. str. PR17 AF2298
AF229879.1 gg_id:24208 Ochrobactrum sp. str. 4FB13 AF2298
FJ859687.2 gg_id:536560 Ochrobactrum pseudogrignonense str. BIHB 340 FJ8596
DQ112028.1 gg_id:130527 Ochrobactrum anthropi str. T1 DQ112
AM084018.1 gg_id:140347 Ochrobactrum sp. str. R-25055 AM084
AM490612.1 gg_id:249691 Ochrobactrum anthropi str. LMG 401 AM490
AJ550273.2 gg_id:90730 Ochrobactrum sp. TK14 AJ5502
DQ295875.1 gg_id:142896 Ochrobactrum sp. str. CDB2 DQ295
EU370421.1 gg_id:271876 Ochrobactrum sp. str. 45C2 EU370
EU352762.1 gg_id:268033 Ochrobactrum tritici str. NK 2.X-2 EU352
DQ133574.1 gg_id:139474 Ochrobactrum sp. str. LJJS1-2 DQ133
EF028152.1 gg_id:179539 Ochrobactrum sp. str. xll-3 EF0281
EU399793.1 gg_id:272191 Ochrobactrum anthropi str. PV2Z2 EU399
AM411061.1 gg_id:175969 Ochrobactrum cytisi str. 43(25)zhy AM411
GQ203110.1 gg_id:585971 Ochrobactrum pseudogrignonense str. 4-4DEP GQ203
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GQ451321.1 gg_id:510928 Fecal Yunnan snub-nosed monkey feces clone J346 GQ451
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FJ493040.1 gg_id:589752 Pseudochrobactrum asaccharolyticum str. 2G-2 FJ4930
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FJ950554.1 gg_id:407459 Pseudochrobactrum asaccharolyticum str. c72 FJ9505
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AB369864.1 gg_id:253214 Pseudochrobactrum glaciale str. KMM 3858 (= Pi 26) AB3698
AM114407.1 gg_id:151281 Ochrobactrum anthropi str. CCUG 34735 AM114
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D25314.1 gg_id:4687 Blastochloris viridis str. ATCC19567 D25314
AM502287.1 gg_id:261580 Blastochloris gulmargensis str. JA248 AM502
AF084509.1 gg_id:200193 Blastochloris viridis str. B9 AF0845
AJ401203.1 gg_id:78031 Blastochloris sp. AT2101 AJ4012
AY117148.1 gg_id:63558 Blastochloris sulfoviridis str. OK3G7 AY1171
AJ012089.1 gg_id:4711 Blastochloris sulfoviridis str. ToP1 AJ0120
AJ548824.1 gg_id:82396 Candidatus Devosia euplotis str. CAMP4.4 AJ5488
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EF433460.1 gg_id:212196 Devosia sp. str. IPL15 EF4334
EU812979.1 gg_id:311842 Microbial structure thermophilic reactors biogas reactor fed corn silage clone TUM-M EU8129
EF540476.1 gg_id:224522 Devosia sp. 6_4V EF5404
FJ807418.1 gg_id:550842 Presence and iron oxidizing landfill leachate impacted groundwater surface water int FJ8074
EU362140.1 gg_id:355188 and Mesquite Flats Site Death Valley dune sand clone 11 EU3621
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EF612399.1 gg_id:255206 historic semiarid lead-zinc mine tailings site soil clone OS-C20 EF6123
GQ396877.1 gg_id:573755 Functional unvegetated perhumid recently-deglaciated soils do not correlate shifts s GQ396
AY512822.1 gg_id:105993 Devosia riboflavina AY5128
DQ279337.1 gg_id:144518 Dynamic during maturation Tuber magnatum Pico ascoma Xanthomonas clone TM1 DQ279
EF067859.1 gg_id:174315 Devosia terrae str. DCY11 EF0678
FN421512.1 gg_id:573013 proteogenomics physiology phyllosphere Arabidopsis thaliana clone 12_F07 FN4215
EU289427.1 gg_id:245814 metagenomic approach biosynthesis ansamycin endophytes or symbionts enriched s EU2894
GQ472390.1 gg_id:544772 Microbial Dongping Lake lake sediment clone 4D3-5 GQ472
FM253613.1 gg_id:333986 Rock gold mine unusually microbial biofilm clone F07-1 FM253

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FN594684.1 gg_id:559356 and arsenic mobilization rock biofilms ancient biofilom gold mine Zloty Stok clone f1- FN594
EF018449.1 gg_id:215447 aspen rhizosphere clone Amb_16S_1074 EF0184
GQ263214.1 gg_id:581749 Structure Simulated Level Waste Site simulated level waste site clone F3_19X GQ263
EU709017.1 gg_id:301593 Devosia sp. str. B2 EU709
FJ560749.1 gg_id:550691 Prosthecomicrobium mishustinii str. 17 FJ5607
EF540486.1 gg_id:227483 Devosia sp. c4M EF5404
AM293857.1 gg_id:237298 Devosia subaequoris str. HST3-14 AM293
GU444083.1 gg_id:516056 Effects amendments on fertility and microbial structure cucumber Cucumis sativus L GU444
EF540440.1 gg_id:224543 oil-shale semi-coke solid waste clone PK_X EF5404
CU927436.1 gg_id:573246 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU927
FM872901.1 gg_id:365288 occupant source house floor dust clone FB04D08 FM872
EF575560.1 gg_id:232353 Devosia sp. str. BD-c194 EF5755
AY921786.1 gg_id:114008 farm soil clone AKYG1666 AY9217
EU344945.1 gg_id:273185 Hepialus gonggaensis guts equalized library larval gut clone Hg5-35 EU3449
GQ264021.1 gg_id:540672 Structure Simulated Level Waste Site simulated level waste site clone WC3_112 GQ264
AJ318163.1 gg_id:25284 Waste-gas biofilter clone BIri24 AJ3181
AY695724.1 gg_id:107598 USA: New Mexico Carlsbad Lechuguilla Cave Rainbow Room clone RR3A4 AY6957
GQ263858.1 gg_id:565555 Structure Simulated Level Waste Site simulated level waste site clone WC2_101 GQ263
EU937979.1 gg_id:332501 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU937
CU918672.1 gg_id:552248 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU918
GQ396880.1 gg_id:560208 Functional unvegetated perhumid recently-deglaciated soils do not correlate shifts s GQ396
DQ814270.1 gg_id:163314 zebrafish digestive tract clone aab66c07 DQ814
EU133360.1 gg_id:257399 tallgrass prarie soil clone FFCH11455 EU133
DQ248307.1 gg_id:142990 carbon tetrachloride contaminated soil clone TF11 DQ248
GQ093782.1 gg_id:453654 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ093
Y14309.1 gg_id:4622 Hyphomicrobium facile str. H-526 Y14309
FM877980.1 gg_id:553425 Microbial iron redox cycling circumneutral-pH groundwater seep Hyphomicrobium c FM877
Y14307.1 gg_id:4627 Hyphomicrobium methylovorum str. DSM 5458 Y14307
AB016812.1 gg_id:4626 Hyphomicrobium methylovorum str. MBIC3196 AB0168
AF235089.1 gg_id:29178 Hyphomicrobium sulfonivorans str. S1; ATCC BAA-113; DSM 13863 AF2350
FJ536921.1 gg_id:594171 Linking performance microbiology biofilters treating dimethyl sulphide presence and FJ5369
Y14305.1 gg_id:4632 Hyphomicrobium zavarzinii str. ZV-622 Y14305
AB252934.1 gg_id:167855 Identification Shibayama lagoon Ishikawa Japan iron-oxidation biofilm clone 225 AB2529
EF019299.1 gg_id:210886 aspen rhizosphere clone Elev_16S_489 EF0192
AF198623.1 gg_id:47454 Hyphomicrobium chloromethanicum str. CM2 AF1986
AF234719.1 gg_id:52859 sludge clone A27b AF2347
AY328655.1 gg_id:88846 drinking water system simulator clone DSSF33 AY3286
Y14302.1 gg_id:4636 Hyphomicrobium vulgare str. IFAM MC-750 ATCC 27500 Y14302
DQ814327.1 gg_id:172883 zebrafish digestive tract clone aab67a09 DQ814

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AY328697.1 gg_id:88935 drinking water system simulator clone DSSF75 AY3286
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AF279787.1 gg_id:26693 Hyphomicrobium sp. str. PMC AF2797
AY328635.1 gg_id:89164 drinking water system simulator clone DSSF12 AY3286
AY328650.1 gg_id:90267 drinking water system simulator clone DSSF27 AY3286
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DQ814766.1 gg_id:172128 zebrafish digestive tract clone aab53a04 DQ814
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GQ264377.1 gg_id:513073 Structure Simulated Level Waste Site simulated level waste site clone WW2_57 GQ264
AY328683.1 gg_id:88276 drinking water system simulator clone DSSF61 AY3286
DQ815221.1 gg_id:168134 zebrafish digestive tract clone aab51g05 DQ815
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GQ281066.1 gg_id:586097 Hyphomicrobium facile str. NBCS23 GQ281
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FJ439830.1 gg_id:550289 Investigation estrogen-degrading bioreactors membrane bioreactor biofilm clone M2 FJ4398
AY328674.1 gg_id:90421 drinking water system simulator clone DSSF52 AY3286
EF125445.1 gg_id:198749 mangrove soil clone MSB-2E9 EF1254
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GU269548.1 gg_id:579387 Pedomicrobium ferrugineum str. ATCC 33119 GU269
GQ339120.1 gg_id:529888 Comprehensive Investigation on Iron Cycling Freshwater Seep ing Microscopy Cultiv GQ339
GQ264219.1 gg_id:585537 Structure Simulated Level Waste Site simulated level waste site clone WW1_b13 GQ264
FJ478816.1 gg_id:573426 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4788
FJ479530.1 gg_id:554597 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4795
AY921654.1 gg_id:113465 farm soil clone AKYG1791 AY9216
FJ478521.1 gg_id:556996 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4785
GU269549.1 gg_id:571598 Pedomicrobium manganicum str. ATCC 33121 GU269
EF019229.1 gg_id:207866 aspen rhizosphere clone Elev_16S_398 EF0192
EU644219.1 gg_id:285061 and Samoylov Island Lena active layer -centred polygon near surface horizon; Rim_T EU6442

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EF153296.1 gg_id:187340 Rhodomicrobium sp. str. ShRmc01 EF1532
EF650482.1 gg_id:238973 Rhodocista sp. str. GP-7 EF6504
D25313.1 gg_id:4680 Rhodoplanes roseus str. 941 D25313
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EF019983.1 gg_id:212484 aspen rhizosphere clone Elev_16S_1370 EF0199
EU133616.1 gg_id:254348 tallgrass prarie soil clone FFCH2026 EU1336
EU335329.1 gg_id:245987 Changes microbial metabolic and along hydrogeochemically variable profile unsatura EU3353
AY921820.1 gg_id:113474 farm soil clone AKYG712 AY9218
AY922092.1 gg_id:114396 farm soil clone AKYH1064 AY9220
EF018397.1 gg_id:208293 aspen rhizosphere clone Amb_16S_1010 EF0183
GQ263521.1 gg_id:534454 Structure Simulated Level Waste Site simulated level waste site clone FW2_47B GQ263
EU335450.1 gg_id:235975 Changes microbial metabolic and along hydrogeochemically variable profile unsatura EU3354
EF020096.1 gg_id:210974 aspen rhizosphere clone Elev_16S_1506 EF0200
GQ500727.1 gg_id:576799 Surveys Biofilms and Sediments Mammoth Cave Karst Aquifers Charon's Cascade sa GQ500
AY622245.1 gg_id:107444 subsurface soil clone B-L17 AY6222
FJ479010.1 gg_id:561693 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4790
FJ625323.1 gg_id:570840 Influence lead contamination on boreal pine forest soil clone HF_M_20 FJ6253
EF019513.1 gg_id:207374 aspen rhizosphere clone Elev_16S_1015 EF0195
EF032737.1 gg_id:179877 cyanobacterial mat in lava cave clone HAVOmat148 EF0327
EF018835.1 gg_id:215423 aspen rhizosphere clone Amb_16S_1316 EF0188
AY234497.1 gg_id:111439 soil isolate Ellin5080 AY2344
EU937846.1 gg_id:315986 Biogeochemistry Iron Oxidation Circumneutral Freshwater Habitat riparian iron oxidi EU9378
AY921720.1 gg_id:113725 farm soil clone AKYH1040 AY9217
FJ807419.1 gg_id:570413 Presence and iron oxidizing landfill leachate impacted groundwater surface water int FJ8074
EF020128.1 gg_id:215872 aspen rhizosphere clone Elev_16S_1545 EF0201
DQ537531.1 gg_id:163571 manganese oxidation soil clone JH-GY22 DQ537
FJ152854.1 gg_id:316335 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX5A_ FJ1528
DQ984560.1 gg_id:237595 Mt. Nan-Jen soil clone IFD_31 DQ984
AY326602.1 gg_id:136172 Amazon soil clone 711-1 AY3266
AJ581617.1 gg_id:93792 soil near uranium mill tailings clone KCM-C-24 AJ5816
FJ479160.1 gg_id:551343 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4791
FJ479575.1 gg_id:585582 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4795
AB286367.1 gg_id:204919 Reducing production and domination comamonadaceae reducing oxygen supply was AB2863
EF018437.1 gg_id:212776 aspen rhizosphere clone Amb_16S_1057 EF0184
AY395414.1 gg_id:105676 pasture soil clone EB1095 AY3954
EF019188.1 gg_id:208963 aspen rhizosphere clone Amb_16S_1857 EF0191
DQ451455.1 gg_id:156816 forest soil clone FAC16 DQ451
EU133294.1 gg_id:251152 tallgrass prarie soil clone FFCH12843 EU1332
AY913382.1 gg_id:113588 forest soil clone DUNssu176 (-7A) (OTU#160) AY9133
AY963313.1 gg_id:151184 structures and correlated ecological factors soils two evergreen broad-leaved forests AY9633

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DQ837247.1 gg_id:244310 Archaeal and pristine National Park Spain Donana coastal aquifer clone 49S1_2B_37 DQ837
FJ478974.1 gg_id:584077 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4789
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AY724055.2 gg_id:111888 soil clone DUNssu006 AY724
AB121773.1 gg_id:98457 Bradyrhizobium sp. str. Shinshu-th2 AB121
AY921846.1 gg_id:114139 farm soil clone AKYG1898 AY921
GQ397068.1 gg_id:575164 Functional unvegetated perhumid recently-deglaciated soils do not correlate shifts s GQ397
EU335327.1 gg_id:250714 Changes microbial metabolic and along hydrogeochemically variable profile unsatura EU335
AJ810382.1 gg_id:159481 Rhizobiales genosp. AB CsBT10/2 AJ8103
EU429497.1 gg_id:544454 producing pigmented colonies aerosol and soil remote glaciated areas (Antarctica Al EU429
EF516064.1 gg_id:218539 grassland soil clone FCPS586 EF516
GU444101.1 gg_id:515709 Effects amendments on fertility and microbial structure cucumber Cucumis sativus L GU444
GQ402598.1 gg_id:566207 Accessing microbial biodiversity and metabolic potential tropical -based approach p GQ402
EF018075.1 gg_id:211128 aspen rhizosphere clone Amb_16S_400 EF018
DQ404666.1 gg_id:152904 contaminated sediment clone 654997 DQ404
EU644207.1 gg_id:282627 and Samoylov Island Lena active layer -centred polygon near surface horizon; Rim_T EU644
GQ264285.1 gg_id:555816 Structure Simulated Level Waste Site simulated level waste site clone WW2_06 GQ264
EF019849.1 gg_id:205156 aspen rhizosphere clone Elev_16S_1206 EF0198
DQ125920.1 gg_id:136171 uranium contaminated soil clone AKAU4191 DQ125
EU669630.1 gg_id:314708 soil clone S22-33 EU669
EF020062.1 gg_id:213736 aspen rhizosphere clone Elev_16S_1469 EF020
AY426439.1 gg_id:101471 full-scale anaerobic UASB bioreactor treating paper mill wastewater clone B14 AY4264
EF018152.1 gg_id:216748 aspen rhizosphere clone Amb_16S_487 EF0181
CU918797.1 gg_id:534993 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU918
EU133549.1 gg_id:253944 tallgrass prarie soil clone FFCH2918 EU133
AB240329.1 gg_id:144888 microbial structure Phragmites rhizosphere biofilm reed bed reactor laboratory clon AB240
GQ264159.1 gg_id:584598 Structure Simulated Level Waste Site simulated level waste site clone WW1_a15 GQ264
GQ263836.1 gg_id:592746 Structure Simulated Level Waste Site simulated level waste site clone WC1_b83 GQ263
EU289423.1 gg_id:256335 metagenomic approach biosynthesis ansamycin endophytes or symbionts enriched s EU289
GQ500841.1 gg_id:546305 Surveys Biofilms and Sediments Mammoth Cave Karst Aquifers Bio-Sep beads conta GQ500
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EU335194.1 gg_id:256477 Changes microbial metabolic and along hydrogeochemically variable profile saturate EU335
GQ093920.1 gg_id:452881 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ093
EU289515.1 gg_id:256361 metagenomic approach biosynthesis ansamycin endophytes or symbionts enriched s EU289
EF520408.1 gg_id:228499 structure lakes: what controls ? acid-impacted lake clone ADK-BTh02-5 EF5204
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EU133623.1 gg_id:254308 tallgrass prarie soil clone FFCH929 EU133
EU881166.1 gg_id:317632 Karst region primary forest soil clone KPF200711-134 EU881
DQ125603.1 gg_id:136238 uranium contaminated soil clone AKAU3608 DQ125
DQ269121.1 gg_id:191415 marine macro-alga surface clone DPC255 DQ269

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AB240517.1 gg_id:141913 microbial structure rhizosphere biofilm Sapporo root-base (80 160 mm) Phragmites S AB240
EF520393.1 gg_id:233793 structure lakes: what controls ? acid-impacted lake clone ADK-BTh02-93 EF5203
AY934488.1 gg_id:138039 Hyphomicrobium sp. str. WG6 AY9344
GQ397055.1 gg_id:583185 Functional unvegetated perhumid recently-deglaciated soils do not correlate shifts s GQ397
AM934937.1 gg_id:572993 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM934
EU491418.1 gg_id:263450 Abundance and microbial life ocean crust seafloor lavas Hawai'i South Point X4 clone EU491
AM935138.1 gg_id:564931 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM935
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DQ413077.1 gg_id:151855 Microbial -dependent and Approaches Anaerobic/Aerobic SBR Reactor EBPR sludge DQ413
EU491437.1 gg_id:274058 Abundance and microbial life ocean crust seafloor lavas Hawai'i South Point X4 clone EU491
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DQ860029.1 gg_id:248682 anchovy intestinal microflora clone F1W DQ860
FM242368.1 gg_id:305774 sequences sediment clone 64 T9d+oil FM242
GQ264526.1 gg_id:589466 Structure Simulated Level Waste Site simulated level waste site clone WW3_80 GQ264
DQ125521.1 gg_id:136194 uranium contaminated soil clone AKAU3482 DQ125
FM253663.1 gg_id:344787 Rock gold mine unusually microbial biofilm clone F01-2 FM253
EU488066.1 gg_id:552142 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU488
EU491718.1 gg_id:262760 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU491
GU362945.1 gg_id:572272 and archaeal biogeochemical cycling genes within Qiongdongnan basin Sea: pilot stu GU362
EU491544.1 gg_id:260534 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR4C EU491
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FM872754.1 gg_id:363198 occupant source house floor dust clone FB01E01 FM872
EU246798.1 gg_id:256471 polyphasic description Pocillopora meandrina Palmyra Atoll Calcinus obscurus abdon EU246
AB453748.1 gg_id:550276 Sulfur-Oxidizing Deep-sea Sponges deep-sea Poecilosclerida sponge clone GM-BSS-B AB453
AB487448.1 gg_id:574086 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNIR0 AB487
EF117253.1 gg_id:178313 Hyphomicrobium sp. str. SLG5B-19 EF1172
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AB099999.1 gg_id:150659 Microbial dead chimney structures systems Mid-Okinawa Trough and Indian Ocean F AB0999
EF018886.1 gg_id:205881 aspen rhizosphere clone Amb_16S_1374 EF0188
EU753663.1 gg_id:353953 modern freshwater stromatolites Ruidera Pools Natural Park Spain dry stromatolite EU753
EF540511.1 gg_id:226508 Devosia sp. 4_C16_46 EF5405
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AB252205.1 gg_id:238917 Methylobacterium brachiatum str. Hojyo2 AB252
EF116592.1 gg_id:190418 Methylobacterium fujisawaense str. TNAU14 EF1165
FJ382176.1 gg_id:574655 Potentially pathogenic and air stem cell transplant summer hospital shower water BM FJ3821
GQ181028.1 gg_id:588577 Methylobacterium fujisawaense str. LNW7 GQ181
EF015477.1 gg_id:189399 Methylobacterium fujisawaense str. MP1 EF0154
EU379265.1 gg_id:266548 Methylobacterium hispanicum str. 2N-2A EU379
EU867320.1 gg_id:581205 Methylobacterium hispanicum str. CCGE2229 EU867
EU379303.1 gg_id:259601 Methylobacterium hispanicum str. 6R-J-4b EU379
AJ888240.1 gg_id:132019 Methylobacterium sp. str. AR25 AJ8882
AY169421.1 gg_id:81189 Methylobacterium fujisawaense AY169
GQ113305.1 gg_id:404095 Topographical and Temporal Human Skin Microbiome skin retroauricular crease clor GQ113
FJ593719.1 gg_id:528281 midgut tropical caterpillars fed on diets egg clone 298EGG8 FJ5937
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AB252201.1 gg_id:237732 Methylobacterium komagatae str. 002-079 AB252
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NC_010725.1 gg_id:289369 Methylobacterium populi str. BJ001 NC_01
GU294326.1 gg_id:574537 Methylobacterium radiotolerans str. 5310 GU294
AB175637.1 gg_id:152031 Methylobacterium organophilum str. DSM 760 AB175
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AB175646.1 gg_id:151826 Methylobacterium thiocyanatum str. DSM 11490 AB175
GQ281068.1 gg_id:543707 Methylobacterium extorquens str. NBCS12 GQ281
D32230.1 gg_id:4776 Methylobacterium zatmanii str. JCM 2819 D3223
AB175647.1 gg_id:150755 Methylobacterium zatmanii str. DSM 5688 AB175
GU294335.1 gg_id:543270 Methylobacterium zatmanii str. 6012 GU294
EF682985.1 gg_id:242745 Eastern Mediterranean atmosphere clone F13_9A_FL EF6829
GQ084916.1 gg_id:374179 Topographical and Temporal Human Skin Microbiome skin manubrium clone nbw26 GQ084
U58018.1 gg_id:4788 Methylobacterium thiocyanatum str. ALL/SCN-P U5801
AY741725.1 gg_id:101778 Methylobacterium sp. str. SKJH-20 AY741
DQ507203.1 gg_id:159608 Methylobacterium sp. str. K611B DQ507

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EF116586.1 gg_id:178333 Methylobacterium thiocyanatum str. TNAU8 EF1165
AY436812.1 gg_id:105470 Methylobacterium sp. str. ST4.9 AY4368
AY616142.1 gg_id:110238 Methylobacterium radiotolerans str. P AY6161
EU528225.1 gg_id:284643 Seasonal Dynamics Mudflat Mouth Major Kentucky Lake Reservoir Tributary sedime EU5282
AB220086.1 gg_id:134057 Methylobacterium sp. str. PB140 AB2200
AB175635.1 gg_id:154997 Methylobacterium lusitanum str. NCIMB 13779 AB1756
GQ002387.1 gg_id:507275 Topographical and Temporal Human Skin Microbiome skin buttock clone nbu176g11 GQ002
AY838487.1 gg_id:109372 fruiting body Pleurotus eryngii clone PE31 AY8384
EU379312.1 gg_id:277255 Methylobacterium fujisawaense str. 7RN-5A EU3793
M95654.1 gg_id:4785 Methylobacterium sp str. PK-1 M9565
EF015479.1 gg_id:196467 Methylobacterium lusitanum str. MP2 EF0154
DQ339624.1 gg_id:146461 Methylobacterium sp. str. Ens25 DQ339
EF116591.1 gg_id:177690 Methylobacterium aminovorans str. TNAU13 EF1165
AM910532.1 gg_id:268349 Methylobacterium aminovorans str. F13 AM910
GQ008625.1 gg_id:501040 Topographical and Temporal Human Skin Microbiome skin antecubital fossa clone n GQ008
NR_029082.1 gg_id:584448 Methylobacterium populi BJ001 str. BJ001; ATCC BAA-705; NCIMB 13946 NR_029
AF198624.1 gg_id:52319 Methylobacterium chloromethanicum str. CM4 AF1986
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AY366072.1 gg_id:93863 Methylobacterium extorquens str. UCM B-3368 AY3660
AF267912.1 gg_id:43152 Methylobacterium extorquens str. DSM 13060 F AF2679
EF116584.1 gg_id:190499 Methylobacterium suomiense str. TNAU4 EF1165
EF116583.1 gg_id:180676 Methylobacterium populi str. TNAU3 EF1165
FJ225120.1 gg_id:332818 Methylobacterium sp. str. DDW-1 FJ2251
DQ223682.1 gg_id:161482 Methylobacterium sp. zf-IVRht11 DQ223
EF426729.1 gg_id:204294 Methylobacterium sp. PMB02 EF4267
AB252208.1 gg_id:253017 Methylobacterium tardum str. RB677 AB2522
AY365229.1 gg_id:93518 Methylobacterium extorquens str. UCM B-3362 AY3652
AB252209.1 gg_id:246889 Methylobacterium gregens str. RB678 AB2522
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AB369041.1 gg_id:555620 and microbial oils produced Asia petroleum crude oil (Qatar Land Light) clone PT-QTL AB3690
EU250227.1 gg_id:249628 acid mine drainage clone GXDC-13 EU2502
AB220102.1 gg_id:138655 Methylobacterium sp. str. PB236 AB2201
FJ192175.1 gg_id:329760 Genetic inventory 'KSC-PHSF' spacecraft assembly clean room during Phoenix missior FJ1921
EF116582.1 gg_id:182127 Methylobacterium populi str. TNAU1 EF1165
AY741717.1 gg_id:105899 Methylobacterium sp. str. SKJH-15 AY7417
D32228.1 gg_id:4786 Methylobacterium rhodesianum str. JCM 2810 D32228
AJ851087.1 gg_id:112210 Methylobacterium variabile str. GR3 AJ8510
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DQ471331.1 gg_id:157883 Methylobacterium jeotgali str. S2R03-9 DQ471
AF409010.1 gg_id:102764 Methylobacterium sp. str. Ellin168 AF409
Z23159.1 gg_id:4797 Methylobacterium sp. str. PC30.44 (pale pink colony type) Z23159
AB252204.1 gg_id:312016 Methylobacterium sp. Hojyo1 AB252
D32229.1 gg_id:4784 Methylobacterium rhodinum str. JCM 2811 D3222
DQ129299.1 gg_id:138274 microarray level air two cities Texas urban aerosol clone AKIW886 DQ129
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AF107461.1 gg_id:32547 Methylosinus pucelana str. MTS AF107
AJ458508.1 gg_id:64197 Methylocystis parvus str. 57 AJ4585
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AJ458467.1 gg_id:64230 Methylocystis sp. str. 495/4 AJ4584
FJ502233.3 gg_id:558957 Methylopila sp. str. JZL-4 FJ5022
DQ404188.1 gg_id:212879 Schlegelia plantiphila str. S1 DQ404
AB007842.1 gg_id:58407 methylotroph str. Hohenstaufen AB007
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AJ458495.1 gg_id:111428 Methylosinus sporium str. KS16a AJ4584
M95664.1 gg_id:4735 Methylosinus sp str. LAC M9566
AJ458481.1 gg_id:64219 Methylosinus trichosporium str. 33/1 AJ4584
AF150804.1 gg_id:4739 Methylosinus trichosporium AF150
L20845.1 gg_id:4736 methanotrophic str. IMV B-3060 L20845
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EF515279.1 gg_id:220495 Electricigen Enrichment MFC upf microbial fuel cell anode clone 22c11 EF5152
CU924338.1 gg_id:560880 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU924
AB127972.1 gg_id:101532 Kaistina koreensis str. Y9 AB127
GQ060654.1 gg_id:467965 Topographical and Temporal Human Skin Microbiome skin glabella clone nbw03g03 GQ060
GQ064493.1 gg_id:418162 Topographical and Temporal Human Skin Microbiome skin axillary vault clone nbw7 GQ064
AJ458504.1 gg_id:64202 Methylocystis sp. str. F10V2a AJ4585
AB095950.1 gg_id:87497 Ancalomicrobium adetum AB095
AJ458500.1 gg_id:64205 Methylocystis sp. str. 10 AJ4585
GQ055294.1 gg_id:371984 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw1113d GQ055
CU925731.1 gg_id:548842 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU925

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AY138237.1 gg_id:107461 human stomach biopsy clone Sto3-1 AY138
AJ458473.1 gg_id:56422 Methylocystis echinoides str. IMET 10491 AJ4584
GQ402774.1 gg_id:537864 Accessing microbial biodiversity and metabolic potential tropical -based approach p GQ402
AM283543.1 gg_id:165785 Methylocystis heyerii str. H2 AM283
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GU208455.1 gg_id:591539 Study on Sediment Dongping Lake sediment clone Fr1-9 GU208
GQ500824.1 gg_id:540718 Surveys Biofilms and Sediments Mammoth Cave Karst Aquifers Bio-Sep beads conta GQ500
GQ500812.1 gg_id:556729 Surveys Biofilms and Sediments Mammoth Cave Karst Aquifers Bio-Sep beads conta GQ500
AM934956.1 gg_id:546522 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM934
FJ755762.1 gg_id:552837 vertical archaeal water and Lake Taihu sediment clone LT-SB-B39 FJ7557
CU923583.1 gg_id:566212 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU923
FJ542854.1 gg_id:530919 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone A FJ5428
CU921271.1 gg_id:550854 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU921
CU926316.1 gg_id:532947 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU926
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AY921960.1 gg_id:113860 farm soil clone AKYG1560 AY921
FM874050.1 gg_id:362109 occupant source house mattress dust clone MB05G07 FM874
EF032656.1 gg_id:182489 structure riverine biofilms river biofilm clone SepB-8 EF0326
EU589298.1 gg_id:272318 China Ningxia rice paddy field soil clone 2_D3 EU589
GQ093788.1 gg_id:382544 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ093
EF590055.2 gg_id:230239 changes structure sediments Songhua River contaminated nitrobenzene -DGGE cloni EF5900
AF370880.1 gg_id:66822 Nordella oligomobilis str. N21 AF3708
DQ984605.1 gg_id:236172 LY oil-contaminated soil clone LYC075 DQ984
AJ582085.1 gg_id:102514 Ahrensia sp. DFL-33 AJ5820
AJ582087.1 gg_id:95153 Ahrensia sp. DFL-44 AJ5820
AJ582086.1 gg_id:94505 Ahrensia sp. DFL-42 AJ5820
AJ582084.1 gg_id:95110 Ahrensia sp. DFL-13 AJ5820
L20802.1 gg_id:5138 Aminobacter ciceronei str. ER2 L20802
DQ196478.1 gg_id:139758 Aminobacter sp. str. MI-p2a DQ196
GQ254285.1 gg_id:544184 Aquamicrobium sp. str. H15 GQ254
FJ167676.1 gg_id:307902 Chelatovorus multitrophus str. LPM-6 FJ1676
EF457242.1 gg_id:237931 Chelatovorus oligotrophus str. LPM4 EF4572
GU111569.1 gg_id:537767 Nitratireductor sp. str. SL014B-25A2 GU111
EF157205.1 gg_id:249985 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF1572
EU564843.1 gg_id:267825 Nitratireductor sp. str. CC-MHSW-5 EU5648
EU281628.1 gg_id:243131 Bacillus sp. B2(2007b) str. B2 EU281
EU024398.2 gg_id:238204 Isolation cyclohexylalkyltetralin degradation sediments marine sediment clone WBI3 EU024

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EU849580.1 gg_id:554748 Mesorhizobium sp. str. CCNWYC132 EU8495
EU410952.1 gg_id:276865 Mesorhizobium temperatum str. CCNWSX0012-2 EU4109
AY195843.1 gg_id:82201 Mesorhizobium mediterraneum str. PECA03 AY1958
EU074171.1 gg_id:253654 Mesorhizobium mediterraneum str. CCBAU 01409 EU0741
AF508208.1 gg_id:102422 Mesorhizobium temperadae str. CCBAU 11018; SDW 018 AF5082
AY725261.1 gg_id:136815 decayed velvetleaf seed clone S1-4-CL6 AY7252
FJ234902.1 gg_id:352327 Microbial structure mesophilic biogas mesophil anaerobic reactors fed corne silage cl FJ2349
AF410897.1 gg_id:20161 Mesorhizobium sp. str. Lo7-12 AF4108
FJ550306.1 gg_id:512978 Mesorhizobium tianshanense str. BIHB 4151 FJ5503
FJ025121.1 gg_id:324436 Mesorhizobium tianshanense str. SEMIA 3012 FJ0251
EU707156.1 gg_id:290008 Mesorhizobium tianshanense str. CCBAU 15528 EU7071
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GQ500629.1 gg_id:558777 Mesorhizobium sp. str. S948 GQ500
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DQ100068.1 gg_id:135924 Mesorhizobium plurifarium str. CCBAU 51276 DQ100
AY922072.1 gg_id:113594 farm soil clone AKYG882 AY9220
FJ025123.1 gg_id:306937 Mesorhizobium amorphae str. SEMIA 849 FJ0251
AF325770.1 gg_id:47902 Rhizobium sp. str. TM-5 AF3257
AY688622.1 gg_id:136093 Mesorhizobium plurifarium str. Sp42 AY6886
DQ859034.1 gg_id:204637 Mesorhizobium plurifarium str. ORS 3324 DQ859
FJ550307.1 gg_id:548722 Mesorhizobium tianshanense str. BIHB 4154 FJ5503
D14514.1 gg_id:5142 Mesorhizobium loti str. ATCC33669 D14514
EU256441.1 gg_id:248289 Mesorhizobium loti str. CCBAU 85068 EU2564
D14511.1 gg_id:5149 Rhizobium sp str. TAL 1148 D14511
GQ088394.1 gg_id:443339 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw: GQ088
AF508207.1 gg_id:102269 Mesorhizobium nortea str. CCBAU 11014; SDW 014 AF5082

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AB166876.1 gg_id:99830 Mesorhizobium sp. str. M01 AB1668
AF041447.1 gg_id:5146 Mesorhizobium tianshanense str. -1BS; USDA 3592 AF0414
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FJ542962.1 gg_id:511099 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone B FJ5429
AJ295079.1 gg_id:74017 Mesorhizobium plurifarium str. ORS1096 (LMG 15298) AJ2950
EU874893.1 gg_id:308046 Mesorhizobium sp. W12 EU8748
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CU915133.1 gg_id:381876 and dynamics adapted microbiota during degradation Tunisian zarzatine enriched se CU915
EU440986.1 gg_id:317496 Nitratireductor aquibiodomus str. PR57-9 EU4409
EU874875.1 gg_id:344328 Mesorhizobium sp. W66 EU8748
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EU143347.2 gg_id:253753 Nitratireductor sp. str. J3 EU1433
DQ659453.1 gg_id:160997 Chelatobacter sp. Pht-3B DQ659
GU447302.1 gg_id:518497 Nitratireductor sp. str. C115 GU447
FJ182044.1 gg_id:313217 Parvibaculum sp. str. P31 FJ1820
DQ015855.1 gg_id:139871 Antarctic lake water clone WLB16-101 DQ015
FJ536887.1 gg_id:563896 Linking performance microbiology biofilters treating dimethyl sulphide presence and FJ5368
AB186821.1 gg_id:101714 polychlorinated dioxin dechlorinating microcosm clone TSAC02 AB1868

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FM242452.1 gg_id:311978 sequences sediment clone 9 T12d+oil FM242
AY387398.1 gg_id:101358 Parvibaculum lavamentivorans str. DS-1; DSM13023; NCIMB13966 AY3873
FN430653.1 gg_id:537152 Parvibaculum sp. str. MBNA2 FN4306
FJ154092.1 gg_id:333292 Phyllobacterium ifriqiyense str. CCNWGS0184 FJ1540
EU169165.1 gg_id:236290 Phyllobacterium myrsinacearum str. B65 EU1691
EU420076.1 gg_id:541971 Phyllobacterium myrsinacearum str. Sp-0 EU4200
AY636000.1 gg_id:104486 Phyllobacterium callistus str. CSC19 AY6360
DQ917820.1 gg_id:219107 Microbial profiles bleached and wild-type Muricea elongata Phyllobacterium clone B DQ917
FJ405362.1 gg_id:367932 Phyllobacterium myrsinacearum str. SPc FJ4053
EU169173.1 gg_id:253647 Phyllobacterium myrsinacearum str. B18 EU1691
AY785319.1 gg_id:144696 Phyllobacterium brassicacearum str. STM 196 AY7853
AJ011330.1 gg_id:5118 Phyllobacterium myrsinacearum HM35 AJ0113
EU928863.1 gg_id:345333 Phyllobacterium sp. str. T1018 EU9288
X98418.1 gg_id:5120 Nanobacterium sanguineum Seralab FBS batch 901045 X98418
AY785329.1 gg_id:142779 Phyllobacterium myrsinacearum str. STM 958 AY7853
EF364390.1 gg_id:225459 Phyllobacterium sp. str. Hue-5 EF3643
AY635999.1 gg_id:107799 Phyllobacterium callistus str. CSC32 AY6359
AY512821.1 gg_id:105844 Phyllobacterium myrsinacearum AY5128
GU257961.1 gg_id:593356 Bartonella elizabethae str. HXG-C15 GU257
AY331575.1 gg_id:91742 Pseudaminobacter sp. str. mp-1 AY3315
D32248.1 gg_id:5112 Thiobacillus sp. str. THI 051 D32248
FJ203076.1 gg_id:337809 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2030
FJ203650.1 gg_id:356370 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2036
GU118765.1 gg_id:542574 Threatened Corals Provide Microbial Habitats clone Mfra_M12 GU118
FJ203080.1 gg_id:338592 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2030
FJ203059.1 gg_id:303869 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2030
FJ203057.1 gg_id:350236 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2030
DQ396147.1 gg_id:155758 deep-sea octacoral clone ctg_NISA060 DQ396
AB099936.1 gg_id:79338 inactive deep-sea hydrothermal vent chimneys clone IheB2-3 AB0999
EU491659.1 gg_id:270912 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU4916
AF358017.1 gg_id:26724 peat soil microcosm clone LO13.3 AF3580
EU491413.1 gg_id:269457 Abundance and microbial life ocean crust seafloor lavas Hawai'i South Point X4 clone EU4914
EF572341.1 gg_id:228614 Coco's Island site 23 marine clone S23_440 EF5723
DQ351777.1 gg_id:147394 Microbial Adherent Sediment Particles Heavy Metal Contaminated North Sea Surfac DQ351
GQ250606.1 gg_id:545163 bulk sea water column Gulf Mexico clone GOM_WB10-41 GQ250
EU246807.1 gg_id:248919 polyphasic description Pocillopora meandrina Palmyra Atoll Calcinus obscurus abdon EU2468
EU488046.1 gg_id:568753 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU4880
FJ478659.1 gg_id:548335 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4786
FM253615.1 gg_id:337543 Rock gold mine unusually microbial biofilm clone F09-1 FM253
EU735674.1 gg_id:343405 archaeal structures oil contaminated and soils China pristine soil Jidong Oilfield clone EU7356
AY913401.1 gg_id:113418 forest soil clone DUNssu195 (-7A) (OTU#173) AY9134

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AY922026.1 gg_id:112779 farm soil clone AKYG492 AY922(
AY913262.1 gg_id:113787 forest soil clone DUNssu042 (-2A) (OTU#176) AY913
GQ263181.1 gg_id:510769 Structure Simulated Level Waste Site simulated level waste site clone F3_139X GQ263
EU881134.1 gg_id:308120 Karst region primary forest soil clone KPF200711-010 EU881
EU236381.1 gg_id:288220 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg91H4 EU236
AY913338.1 gg_id:114183 forest soil clone DUNssu131 (+7A) (OTU#177) AY913
EF019626.1 gg_id:215375 aspen rhizosphere clone Elev_16S_770 EF0196
EU488067.1 gg_id:548592 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU488(
FJ152813.1 gg_id:308393 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX5A_ FJ1528
EU386115.1 gg_id:270010 Biogeographical and microbes sediments on subseafloor sediment South China Sea c EU386
GQ262947.1 gg_id:549426 Structure Simulated Level Waste Site simulated level waste site clone F1_85X GQ262
AJ633940.1 gg_id:114727 host's accessory nidamental gland clone 52ANG2 AJ6339
EU386026.1 gg_id:272493 Biogeographical and microbes sediments on subseafloor sediment South China Sea c EU386(
AY913351.1 gg_id:112770 forest soil clone DUNssu144 (+7A) (OTU#173) AY913
FJ479146.1 gg_id:588630 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4791
EU491428.1 gg_id:260972 Abundance and microbial life ocean crust seafloor lavas Hawai'i South Point X4 clone EU491
DQ070827.1 gg_id:147965 Prokaryotic Structures Seafloor Basalts Varying Age East Pacific Rise and Ridge basal DQ070
GU369888.1 gg_id:529803 and mats vents Arc white filamentous microbial mat photic shal hydrothermal vent GU369
AB491708.1 gg_id:559688 Microbial control lamina formation Crystal geyser central Utah travertine clone CG1(AB491
DQ814512.1 gg_id:161858 zebrafish digestive tract clone aaa30b10 DQ814
GQ262857.1 gg_id:567747 Structure Simulated Level Waste Site simulated level waste site clone F1_33X GQ262
EU491525.1 gg_id:271177 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR4(EU491
AY162106.1 gg_id:150582 Cultivating Sargasso Sea clone GMD21A06 AY162
AY913263.1 gg_id:113892 forest soil clone DUNssu043 (-2A) (OTU#173) AY913
GQ262821.1 gg_id:534554 Structure Simulated Level Waste Site simulated level waste site clone F1_12X GQ262
EU735621.1 gg_id:339556 archaeal structures and pristine soils China oil contaminated soil Jidong Oilfield clone EU735
EU799979.1 gg_id:351580 It's all ranking aquatic Newport Harbour RI clone 1C227658 EU799
EU491596.1 gg_id:264919 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU491
FJ745115.1 gg_id:575156 Transporter genes expressed coastal response dissolved organic carbon surface wate FJ7451
EU385898.1 gg_id:259577 Stratified microbes sediments Core MD05-2902 subseafloor sediment South China S EU385
EU181478.1 gg_id:245815 Prokaryotic enhanced organic matter remineralization continental margin marine se EU181
AF406520.1 gg_id:106379 complex marine community d at depth 200 m North Aegean Sea (Station 3) clone AE AF406
GQ262936.1 gg_id:575894 Structure Simulated Level Waste Site simulated level waste site clone F1_79X GQ262
AM936572.1 gg_id:547134 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM936
GU369924.1 gg_id:518338 and mats vents Arc white microbial film on pebble aphotic zone shal hydrothermal v GU369
AM936135.1 gg_id:553156 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM936
AM712634.1 gg_id:257971 Anderseniella baltica str. BA141 AM712
GU220743.1 gg_id:571033 Microbial mineralization and involved microbial Basin field temperature hydrotherm GU220
FJ152698.1 gg_id:346361 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX1A_ FJ1526
DQ334628.1 gg_id:176463 Impact metals on sediments heavy metal polluted marine sediment clone HB2-33-2 DQ334

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EF208711.1 gg_id:201560 Vertical and horizontal structure microbial sandy carbonate sediment clone Cl75cm.2 EF2087
EU881167.1 gg_id:342537 Karst region primary forest soil clone KPF200711-032 EU881
GQ347790.1 gg_id:592876 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
AF382114.1 gg_id:79210 bacterioplankton clone ZA3318c AF382
FJ203577.1 gg_id:322591 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2035
GU118597.1 gg_id:581316 Threatened Corals Provide Microbial Habitats clone Mfav_F16 GU118
FJ152626.1 gg_id:304121 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX1A_ FJ1526
FJ024326.1 gg_id:358115 Prokaryotic and their role biogeochemical cycling marine basalts basalt Juan de Fuca I FJ0243
DQ395471.1 gg_id:156247 deep-sea octacoral clone ctg_CGOAA69 DQ395
FJ203565.1 gg_id:315490 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2035
EU491292.1 gg_id:276298 Abundance and microbial life ocean crust seafloor lavas Loi'hi Seamount Pisces Peak EU491
GQ848473.1 gg_id:538386 Microbial sediments vents along East Lau Spreading Centre deep sea hydrothermal GQ848
FJ152991.1 gg_id:340224 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX4C FJ1529
GQ009791.1 gg_id:499876 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw77 GQ009
DQ104986.1 gg_id:137534 Mesorhizobium sp. str. JS 12 DQ104
EU332813.1 gg_id:360547 FACS assist identification TFO present aerobic enhanced biological phosphorus remo EU3328
FJ167442.1 gg_id:342632 microbiology granular sludge under autotrophic mixotrophic and heterotrophic denit FJ1674
AB374372.1 gg_id:259747 sequences Weissenstein (white rock) / few millimeters be surface rock called 'Weiss AB3743
FJ203570.1 gg_id:316649 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2035
AB246806.1 gg_id:146335 Mesorhizobium sp. str. Q-2 AB2468
DQ001642.1 gg_id:158376 Microbial seeps asphalt seep clone E101-27 DQ001
EF471634.1 gg_id:269321 Fast Screening and Estuarine Operon Libraries whole surface water Chesapeake Bay EF4716
AF234721.1 gg_id:32754 sludge clone A7 AF2347
AJ864462.1 gg_id:109471 Mesorhizobium thiogangeticum str. SJT AJ8644
FJ674841.1 gg_id:529939 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6748
CU922361.1 gg_id:547097 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU922
AY598817.1 gg_id:105079 Hoeflea marina str. LMG 128 AY5988
NR_025411.1 gg_id:547162 Mesorhizobium chacoense str. PR5; DSM 17287 NR_02
GU118307.1 gg_id:538811 Threatened Corals Provide Microbial Habitats clone Dstr_L15 GU118
GU118284.1 gg_id:589090 Threatened Corals Provide Microbial Habitats clone Dstr_M18 GU118
AF280865.1 gg_id:72907 bioreactor clone mle1-44 AF2808
AB167198.1 gg_id:107911 Rhizobium sp. str. c19 AB167
AF150703.1 gg_id:24111 Elbe River snow isolate Iso26 Iso26_1389 AF1507
FJ670878.1 gg_id:524118 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6708
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FJ152680.1 gg_id:332848 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX1A_ FJ1526
FN436146.1 gg_id:554537 Molekularbiologische Charakterisierung der bakteriellen Populationsdynamik eines t FN436
AY942778.1 gg_id:145164 but species richness marine sponge Cymbastela concentrica clone Cc173 AY9427
AM934887.1 gg_id:573454 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM934
AY690674.1 gg_id:99235 Mesorhizobium sp. str. GC08 AY6906

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FJ671077.1 gg_id:519193 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6710
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AJ318199.1 gg_id:73511 Waste-gas biofilter clone BIwii1 AJ3181
FJ542875.1 gg_id:521776 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone A FJ5428
FJ203541.1 gg_id:316301 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata I FJ2035
GQ263841.1 gg_id:548137 Structure Simulated Level Waste Site simulated level waste site clone WC1_b89 GQ263
EU287328.1 gg_id:243907 Pacific arctic surface sediment clone S26-28 EU2873
FJ671628.1 gg_id:529153 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6716
FJ675300.1 gg_id:527154 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6753
DQ860072.1 gg_id:170119 Symbiotic aerobic anoxygenic phototrophic colonial tunicate clone L9 DQ860
AY307924.1 gg_id:90640 Aminobacter sp. str. COX AY3079
EU236387.1 gg_id:287433 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg92A11 EU2363
EU223941.1 gg_id:246412 -RFLP microbes Qinghai-Tin rail soil clone 56 EU2239
FJ671907.1 gg_id:514109 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6719
DQ814869.1 gg_id:162010 zebrafish digestive tract clone aab54f12 DQ814
GU454867.1 gg_id:523337 Understanding short-chain fatty acids accumulation enhanced alkaline digestion: kir GU454
DQ364238.2 gg_id:157010 Rhizobium sp. str. WH2K DQ364
FJ542989.1 gg_id:528720 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone B FJ5429
Y15403.1 gg_id:5113 Aquamicrobium defluvii str. DSM 11603 Y15403
DQ860063.1 gg_id:167582 Symbiotic aerobic anoxygenic phototrophic colonial tunicate clone C7 DQ860
FJ674998.1 gg_id:518661 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6749
FJ907162.1 gg_id:552671 Aminobacter sp. str. Sokolova FJ9071
FJ542969.1 gg_id:531912 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone B FJ5429
AB257631.1 gg_id:267976 Microbial few millimeters be surface dolomite rock 'Val Piora' Alps 1965 meters abo AB2576
AY515416.1 gg_id:99884 bulk water German Wadden Sea May 1999 clone GWS-BW-H52M AY5154
AB257653.1 gg_id:273074 Microbial few millimeters be surface dolomite rock 'Val Piora' Alps 1965 meters abo AB2576
EU404018.1 gg_id:306709 Microbial septic tank effluent and soil biozone infiltrative surface clean water clone (EU4040
EU769196.1 gg_id:329286 velvetleaf seed clone S4-6-CL4 EU7691
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D14503.1 gg_id:5409 Agrobacterium rubi str. IFO13261 D14503
X67228.1 gg_id:5430 Agrobacterium rubi str. LMG 156 X67228
D12787.1 gg_id:5414 Agrobacterium rubi str. IAM 13569 D12787
AJ389912.1 gg_id:33557 Agrobacterium vitis str. CFBP2736 AJ3899
DQ466575.1 gg_id:156733 Agrobacterium tumefaciens str. B8LH DQ466
FJ493020.1 gg_id:556421 Agrobacterium tumefaciens str. T0H5 FJ4930
EU295450.1 gg_id:241317 Agrobacterium sp. str. LKXJ-3 EU2954
AB208737.1 gg_id:115126 Agrobacterium sp. AEOC022 AB2087
DQ803328.1 gg_id:190446 human fecal clone RL311_aam24b01 DQ803
EU652866.1 gg_id:280086 Agrobacterium sp. str. ZY068 EU6528

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DQ119110.1 gg_id:134359 Zoogloea ramigera DQ119
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AM157353.1 gg_id:142761 Agrobacterium tumefaciens AM157
DQ278905.1 gg_id:143494 Agrobacterium sp. str. 24DA-1 DQ278
EU022589.1 gg_id:330237 Agrobacterium tumefaciens str. KBA3RM1 EU0225
EF025327.1 gg_id:184622 Agrobacterium tumefaciens str. SREPS2 EF0253
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DQ288946.1 gg_id:144288 Agrobacterium tumefaciens TG32 DQ288
GQ258493.1 gg_id:561134 Rhizobium galegae str. Ca2 GQ258
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EU295451.1 gg_id:242798 Agrobacterium sp. str. LKXJ-2 EU2954
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FN640463.1 gg_id:514145 Agrobacterium sp. str. ZX09 FN6404
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AY626383.1 gg_id:99992 Agrobacterium tumefaciens str. ICMP 369 AY6263
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AJ295683.1 gg_id:26338 Agrobacterium tumefaciens str. IrT-JG-6 AJ2956
EU155875.1 gg_id:244531 Agrobacterium tumefaciens str. D6 EU1558
X73041.1 gg_id:5266 Blastobacter aggregatus str. IFAM 1003 X73041
GQ267695.1 gg_id:562519 Agrobacterium tumefaciens str. CCNWSX539 GQ267
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AY576874.1 gg_id:101430 Agrobacterium tumefaciens str. BY6 AY5768
AB535700.1 gg_id:545894 Agrobacterium tumefaciens str. Pl.2-1 AB5357
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AJ306769.1 gg_id:66930 DCP-dechlorinating consortium clone SHA-91 AJ3067
DQ860031.1 gg_id:250206 anchovy intestinal microflora clone F1Y DQ860
AY048887.1 gg_id:33043 rhizosphere clone Q3-24C12 AY0488
AF239848.2 gg_id:47953 Bradyrhizobium japonicum str. PRY65 AF2398
DQ288890.1 gg_id:144720 Ensifer adhaerens DQ288
EU373406.1 gg_id:261377 Agrobacterium tumefaciens str. TPR09 EU3734
AY626382.1 gg_id:99841 Agrobacterium larrymoorei str. ICMP 15237 AY6263
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X68390.2 gg_id:5210 Sinorhizobium saheli str. LMG 7837
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EU399924.1 gg_id:264788 Rhizobium tropici str. CAF333 EU3999
DQ422964.1 gg_id:155498 Sinorhizobium sp. str. Lv6.1Se DQ422
AB487888.1 gg_id:554002 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSSUR0 AB4878
EU420078.1 gg_id:563426 Agrobacterium rhizogenes str. Cy-0 EU4200
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GQ015367.1 gg_id:494299 Topographical and Temporal Human Skin Microbiome skin palm clone nbw767e11c1 GQ015
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EU694389.1 gg_id:299472 Rhizobium huautlense str. MI-7 EU6943
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D14515.1 gg_id:5330 Rhizobium sp str. OK-50 D14515
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GQ287633.1 gg_id:587508 Rhizobium tropici str. Sap110 GQ287
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EU335452.1 gg_id:243842 Changes microbial metabolic and along hydrogeochemically variable profile unsatura EU3354
EU399916.1 gg_id:269266 Rhizobium tropici str. CAF226 EU3999
FJ405384.1 gg_id:358164 Rhizobium tropici str. CAF446 FJ4053
U89832.1 gg_id:5277 Rhizobium tropici str. CIAT 899; USDA 9030 U89832
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FJ405376.1 gg_id:369125 Rhizobium tropici str. CAF344 FJ4053
EU074168.1 gg_id:255982 Rhizobium galegae str. CCBAU 01393 EU0741
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EU488741.1 gg_id:321329 Rhizobium tropici str. CFN 299 EU4887
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AY509217.1 gg_id:99135 Sinorhizobium fredii str. S150 AY5092
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EF364372.1 gg_id:225279 Rhizobium sp. str. Alm-1 EF3643
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EU488749.1 gg_id:336323 Rhizobium tropici str. 233 EU4887
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U86344.1 gg_id:5267 Rhizobium giardinii str. H152 U86344
AM697248.1 gg_id:221087 sequencing libraries indoor dust clone BF0001C089 AM697
Z94803.1 gg_id:5364 Rhizobium genosp. R str. BDV5365 Z94803
GQ181047.1 gg_id:511579 Rhizobium lusitanum str. BLN7 GQ181
EF157275.1 gg_id:243829 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF1572
AY034026.1 gg_id:31634 Rhizobium loessense str. CCBAU 7190B AY0340
AY117667.1 gg_id:63509 Rhizobium tropici str. PRF77 AY1176
EU399920.1 gg_id:264590 Rhizobium tropici str. CAF279 EU3999
CU927615.1 gg_id:567667 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU927
EU748922.1 gg_id:321970 Rhizobium sp. str. ML102 EU7489
EU488756.1 gg_id:313755 Rhizobium tropici str. H52 EU4887
FJ025129.1 gg_id:315875 Rhizobium sp. str. SEMIA 6411 FJ0251
AY064412.2 gg_id:76783 Rhizobium sp. str. DCP2 AY0644
AB233492.1 gg_id:205713 Amorphomonas oryzae str. B26 AB2334
EU867317.1 gg_id:586260 Rhizobium sp. str. CCGE2052 EU8673
U29387.1 gg_id:5389 Rhizobium sp str. RCR 3618D; USDA 2947 U29387
EU256433.1 gg_id:255837 Rhizobium cellulosilyticus str. CCBAU 85045 EU2564
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DQ984537.1 gg_id:256496 FD oil-contaminated soil clone FD_52 DQ984
AF417562.1 gg_id:24266 Rhizobium leguminosarum str. DASA23002 AF4175
AF510386.1 gg_id:87585 Rhizobium sp. str. vgs-11 AF5103
AB233494.1 gg_id:212299 Amorphomonas oryzae str. B47 AB2334
AF417563.1 gg_id:47187 Rhizobium leguminosarum str. DASA23111 AF4175
EU410948.1 gg_id:267876 Rhizobium giardinii str. CCNWGS0022 EU4109
AY117623.1 gg_id:63522 Rhizobium tropici str. PRF34 AY1176
FJ405382.1 gg_id:361567 Rhizobium tropici str. CAF443 FJ4053
Z79620.1 gg_id:5452 Rhizobium galegae str. NBIMTC 2246 Z79620
AY904747.1 gg_id:111491 Agrobacterium rhizogenes str. SEMIA 6070 AY9047
DQ303318.1 gg_id:150175 Dynamic during maturation Tuber magnatum Pico ascoma Rhizobium clone TM4_28 DQ303
EU600086.1 gg_id:276672 Pueraria montana var. lobata EU6000
AM697022.1 gg_id:217449 sequencing libraries indoor dust clone BF0001B048 AM697
EF522124.1 gg_id:220384 Agrobacterium rhizogenes str. CU10 EF5221
D11344.1 gg_id:5282 Rhizobium tropici str. IFO15247 D11344
GQ115602.1 gg_id:459244 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw688b1 GQ115
AF260298.2 gg_id:77278 Rhizobium tropici str. PRF35 AF2602
FJ405375.1 gg_id:359783 Rhizobium tropici str. CAF341 FJ4053
EU488750.1 gg_id:322227 Rhizobium giardinii str. H152 EU4887
AF510382.1 gg_id:87256 Rhizobium sp. str. Kusp-1 AF5103
EU399714.1 gg_id:274169 Rhizobium tropici str. CCBAU 83795 EU3997

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AY316683.1 gg_id:200464 Reichenowia parasiticae	AY316
EF173321.1 gg_id:192524 Sinorhizobium sp. MKI7	EF1733
FJ517717.1 gg_id:560086 Disturbing epithelial homeostasis metazoan Hydra leads drastic changes microbiota e	FJ5177
EU399935.1 gg_id:260090 Rhizobium tropici str. CAF438	EU399
D12788.1 gg_id:5289 Agrobacterium rhizogenes str. IAM 13570	D12788
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EU723213.1 gg_id:334997 Shinella sp. str. NP526a	EU723
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EU346730.1 gg_id:267822 Shinella zoogloeoides str. BC026	EU346
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GQ009804.1 gg_id:499861 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw77	GQ009
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AB285481.1 gg_id:232448 Sulfolanivorax yambarensis str. MS4	AB2854
FJ154090.1 gg_id:351849 Shinella kummerowiae str. CCNWGS0228	FJ1540
EF437254.1 gg_id:249079 Rhizobium sp. str. RZ6	EF4372
FN436071.1 gg_id:573545 Molekularbiologische Charakterisierung der bakteriellen Populationsdynamik eines t	FN4360
FN555404.1 gg_id:511375 Sinorhizobium sp. I-Bh25-4	FN5554
D14255.1 gg_id:5205 Zoogloea ramigera str. IAM12669	D14255
DQ984584.1 gg_id:256527 Mt. Nan-Jen litterfall clone IYF06	DQ984
DQ096644.1 gg_id:132981 Rhizobium sp. PIP4 str. PIP4; JCM 12219	DQ096
EU888002.1 gg_id:349041 Microbial structure hydrolytic stage two-stage anaerobic digestion grass silage leach	EU8880
DQ814239.1 gg_id:162359 zebrafish digestive tract clone aab65g10	DQ814
X74915.1 gg_id:5207 Shinella zoogloeoides str. ATCC 19623	X74915
AM181744.1 gg_id:160209 Sinorhizobium arboris str. LMG 14919	AM181
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AB535696.1 gg_id:557105 Sinorhizobium medicae str. Cl.3-2 AB5356
EU379949.1 gg_id:261990 Sinorhizobium meliloti str. CCBAU 83575 EU3799
AY904727.1 gg_id:111751 Sinorhizobium meliloti str. SEMIA 134 AY9047
EU256452.1 gg_id:244861 Sinorhizobium meliloti str. CCBAU 85092 EU2564
AF533685.1 gg_id:61015 Sinorhizobium meliloti str. WSM1702 AF5336
AM946567.1 gg_id:343932 Ensifer xericitae str. ORS 1401 AM946
GU129566.1 gg_id:594144 Sinorhizobium meliloti str. Qtx-8-1 GU129
AB535703.1 gg_id:545858 Sinorhizobium medicae str. Pl.4-2 AB5357
AB015420.1 gg_id:34931 Sinorhizobium indiaensis str. Ra-3 AB0154
GQ452960.1 gg_id:572099 Sinorhizobium sp. str. DP10 GQ452
EU399794.1 gg_id:275434 Sinorhizobium meliloti str. SaTO3 EU3997
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EF070136.1 gg_id:182386 Sinorhizobium meliloti str. CCBAU 05137 EF0701
AB017203.1 gg_id:50238 Prosthecomicrobium pneumaticum str. MBIC3489 AB017
EU488283.1 gg_id:551681 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU488
FJ793198.1 gg_id:564313 Rhizobium sp. str. CAAS-2R FJ7931
CU920781.1 gg_id:563220 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU920
EU488449.1 gg_id:570051 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU4884
EU488300.1 gg_id:559383 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU4883
EU488378.1 gg_id:565999 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU4883
EU488368.1 gg_id:542279 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU4883
EU488131.1 gg_id:534712 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU4881
AB232339.2 gg_id:155467 Bartonella sp. str. TFC2 AB2323
EU488467.1 gg_id:553301 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU4884
EF540509.1 gg_id:219954 Rhizobium sp. 4_C16_39 EF5405
EF516927.1 gg_id:218154 grassland soil clone FCPP541 EF5169
EU488260.1 gg_id:547565 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU488
AY133103.1 gg_id:56163 TCE-contaminated site clone ccspost2147 AY1331
EU488463.1 gg_id:571469 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU4884
EU488371.1 gg_id:534308 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU4883
EU817493.1 gg_id:326733 Rhizobium sp. str. XJ-L72 EU8174
EF090606.1 gg_id:197918 Agrobacterium sp. str. Mei-QS6 EF0906
EU488459.1 gg_id:565076 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU4884
EU488190.1 gg_id:543717 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU4881
EU488204.1 gg_id:553575 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU488
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EF608175.3 gg_id:535229 Pannonibacter sp. str. HT23 EF6081
AM990762.1 gg_id:279133 Pseudovibrio sp. MOLA 539 AM990
FJ169215.1 gg_id:342767 microbial and polyketide biosynthetic potential palmerolide-producing tunicate Synoi FJ1692
FJ169195.1 gg_id:321382 microbial and polyketide biosynthetic potential palmerolide-producing tunicate Synoi FJ1691
EU726271.1 gg_id:342195 Stappia sp. str. B106 EU726
EU581698.1 gg_id:275806 Rhodobacteraceae sp. MEBiC05064T marine sponge Geomun-Island South Korea EU581
DQ985041.1 gg_id:193166 Halomonas sp. str. JL1044 DQ985
EF101502.1 gg_id:194356 Stappia meyerae str. GA15 EF1015
EU167953.1 gg_id:269625 Stappia sp. str. 16-23 EU1679
AY162083.1 gg_id:135498 Cultivating Sargasso Sea clone GMD37F4 AY1620
FJ202909.1 gg_id:325677 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2029
AY628423.1 gg_id:99161 Stappia marina str. mano18 AY6284
D85832.1 gg_id:111501 Roseibium denhamense str. OCh 254 D85832
AY588950.1 gg_id:122223 Pacific Ocean deep sea sediment clone DE6.8 AY5889
EU168019.1 gg_id:269506 Stappia sp. str. 23-28 EU1680
EU999013.1 gg_id:304551 Predominance Methanolobus and Methanoculleus spp. Archaeal Saline Gas Field Foı EU9990
DQ270709.1 gg_id:144694 Halomonas sp. B-3046 DQ270
GU199002.1 gg_id:593115 Mycoplana sp. str. G110 GU199
GU322907.1 gg_id:524664 Labrenzia sp. str. YC6927 GU322
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DQ985044.1 gg_id:196145 Rhodopseudomonas sp. str. JL1015 DQ985
AM696707.1 gg_id:224191 Rhodobium marinum str. JA211 AM696
GU370102.1 gg_id:521541 Rhodopseudomonas sp. str. P530(9) GU370
EU445271.1 gg_id:272585 Rhodobium pfennigii str. AR 2102 EU445
AM180706.1 gg_id:161325 Rhodobium gokurnum str. JA 173 AM180
DQ532207.1 gg_id:166078 Geographical Variation Microbial Burden and Spacecraft Assembly Johnson Space Ce DQ532
AB374392.1 gg_id:269733 sequences Weissenstein (white rock) / few millimeters be surface rock called 'Weiss AB3743
AB257630.1 gg_id:273081 Microbial few millimeters be surface dolomite rock 'Val Piora' Alps 1965 meters abov AB2576
EU133571.1 gg_id:245274 tallgrass prarie soil clone FFCH4802 EU1335
FJ152795.1 gg_id:310212 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX5A_ FJ1527
DQ451507.1 gg_id:156833 forest soil clone FAC68 DQ451
AY862042.1 gg_id:112593 Washburn Spring 1 community clone WB1B012 AY8620
GQ093827.1 gg_id:452509 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ093
FJ152772.1 gg_id:315470 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX5A_ FJ1527
AF050531.1 gg_id:4714 contaminated aquifer clone WCHB1-55 AF0505
GQ402732.1 gg_id:538998 Accessing microbial biodiversity and metabolic potential tropical -based approach pı GQ402
EU104345.1 gg_id:338628 Evidence Comamonadaceae determinants settling performance activated sludge clo EU1043

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EF516292.1 gg_id:222685 grassland soil clone FCPT613 EF5162
EF465533.1 gg_id:225699 Pseudoxanthobacter soli str. CC-CC4 EF4655
GQ263890.1 gg_id:510008 Structure Simulated Level Waste Site simulated level waste site clone WC2_141 GQ263
EU881257.1 gg_id:332410 (long-term grazing and burned annual) Karst region serious disturbance field soil clor EU881
DQ984620.1 gg_id:257087 YK oil-contaminated soil clone YK_31 DQ984
GQ262949.1 gg_id:538739 Structure Simulated Level Waste Site simulated level waste site clone F1_86X GQ262
AJ229208.1 gg_id:25697 anoxic bulk soil flooded rice microcosm clone BSV56 clone AJ2292
AM162440.1 gg_id:155413 and situ identification acidic sphagnum peat bog clone B83 AM162
U87772.1 gg_id:4760 Afipia genosp. 6 str. G8665 U8777
EF020321.1 gg_id:207430 aspen rhizosphere clone Elev_16S_1897 EF0203
X97077.1 gg_id:4765 peat bog clone TM19 X97077
FJ380167.1 gg_id:360221 ornithogenic Ross Sea region Antarctica AB1 soil clone 13/1/8F FJ3801
AF358012.1 gg_id:29077 peat soil microcosm clone LO13.2 AF358
DQ129631.1 gg_id:137916 microarray level air two cities Texas urban aerosol clone AKIW856 DQ129
DQ823159.1 gg_id:162179 Techniques Microbial Studies Oregon Caves National Monument Quad 3 off trail clo DQ823
GQ350414.1 gg_id:586170 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
EF516212.1 gg_id:225488 grassland soil clone FCPP499 EF5162
FJ152802.1 gg_id:308948 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX5A_ FJ1528
EU881153.1 gg_id:310925 Karst region primary forest soil clone KPF200711-196 EU881
GQ402664.1 gg_id:548721 Accessing microbial biodiversity and metabolic potential tropical -based approach p GQ402
AM936498.1 gg_id:559544 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM936
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GQ402711.1 gg_id:546022 Accessing microbial biodiversity and metabolic potential tropical -based approach p GQ402
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GQ097820.1 gg_id:417366 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw397h0 GQ097
AM935446.1 gg_id:547229 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM935
FM253652.1 gg_id:332834 Rock gold mine unusually microbial biofilm clone D06-2 FM253
EU861875.1 gg_id:338761 effects chronic fertilization on alpine tundra microbial implications carbon and cyclin EU861
CU925666.1 gg_id:553033 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU925
CU926879.1 gg_id:567435 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU926
DQ813987.1 gg_id:162431 zebrafish digestive tract clone aab58b10 DQ813
AY913367.1 gg_id:137744 forest soil clone DUNssu160 (+7B) (OTU#170) AY9133
CU923391.1 gg_id:570766 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU923
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FJ658644.1 gg_id:555885 Metalworking Fluids and Aerosols Metallurgy Plant metalworking aerosol clone Sum FJ6586
NR_025428.1 gg_id:548469 Chelatococcus asaccharovorans str. TE2 NR_02
GQ848214.1 gg_id:552413 and Archaeal Xiarinur Soda Lake Water-sediment Interface water-sediment interfac GQ848
FJ592562.1 gg_id:574090 Fumarole-supported islands biodiversity within hyperarid -elevation landscape on Pu FJ5925
AY792286.1 gg_id:124699 Humic Lake clone CrystalBog021H4 AY792

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FJ542954.1 gg_id:513048 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone B FJ5429
FJ624896.1 gg_id:544990 Influence lead contamination on boreal pine forest soil clone HF_C_23 FJ6248
GQ065523.1 gg_id:452998 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw89d08(GQ065
AY212626.1 gg_id:106732 water 20 m upstream manure clone 175up AY2126
FM253660.1 gg_id:338715 Rock gold mine unusually microbial biofilm clone E09-2 FM253
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AB240233.1 gg_id:144675 microbial structure rhizosphere Phragmites bulk soil reed bed reactor laboratory clo AB240
EU617827.1 gg_id:344447 Seasonal variation microbial Yel Sea sediment clone D13S-4 EU6178
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DQ230971.1 gg_id:141437 subsurface water clone DR938CH110701SACH95 DQ230
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GQ263601.1 gg_id:589565 Structure Simulated Level Waste Site simulated level waste site clone FW3_155C GQ263
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FJ477551.1 gg_id:509105 gut symbionts are tightly linked herbivory ants whole extracted single ant clone 83f FJ4775
FJ416098.1 gg_id:368583 sediments sediment station NEC5 Northern Bering Sea clone 061M64 FJ4160
FJ430076.1 gg_id:360023 Rhizobium sp. str. CCNWGS114 FJ4300
DQ454123.3 gg_id:229692 Rhizobium sp. str. J3-A127 DQ454
EF580937.1 gg_id:277780 Survey Microbial Freshwater Calcareous Mats freshwater calcareous mat clone CM4 EF5809
DQ113411.1 gg_id:143364 microflora ants genus Bartonella Tetraponera attenuata RT13 clone DQ113
GU118131.1 gg_id:535654 Threatened Corals Provide Microbial Habitats clone Apal_F11 GU118
FJ477662.1 gg_id:591010 gut symbionts are tightly linked herbivory ants whole extracted single ant clone UAB- FJ4776
EU056822.1 gg_id:354679 Rhizobium oryzae str. Alt 501 EU0568
DQ269119.1 gg_id:197574 marine macro-alga surface clone DPC247 DQ269
GQ199732.1 gg_id:591717 Mycoplana sp. str. 210_30 GQ199

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EU290679.1 gg_id:254474 microbial structure biogeochemical contrasting shelf sediments Namibian upwelling EU290
EF125460.1 gg_id:176046 mangrove soil clone MSB-2G6 EF1254
EU022307.1 gg_id:256730 Mycoplana dimorpha str. DSM 7138 EU0223
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AJ318143.1 gg_id:46411 Waste-gas biofilter clone BIhii25 AJ3181
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FJ712592.1 gg_id:517638 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) k FJ7125
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FJ193930.1 gg_id:340714 Genetic inventory 'KSC-Astrotek' spacecraft assembly clean room during DAWN missi FJ1939
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AF359524.1 gg_id:38898 Alexandrium lusitanicum NEPCC 253 AF3595
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EF516470.1 gg_id:224138 grassland soil clone FCPO739 EF5164
AB271044.1 gg_id:166471 Labrys ginsengisoli str. Gsoil 106 AB2710
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FM202386.1 gg_id:570096 Hirschia sp. str. GSW-2 FM202
AJ227814.1 gg_id:5865 Hyphomonas jannaschiana str. ATCC 33883 (T) AJ2278
CP000158.1 gg_id:162510 Hyphomonas neptunium str. ATCC 15444 CP0001
AF082795.1 gg_id:5870 Hyphomonas rosenbergii str. VP-6 AF0827
EF540362.1 gg_id:217590 oil-shale semi-coke solid waste clone 07B_MJK EF5403
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DQ160243.2 gg_id:139361 Hyphomonas taiwanensis str. HYP1 DQ160
NR_024938.1 gg_id:572877 Hyphomonas johnsonii str. MHS-2; ATCC 43964 NR_02
AF082797.1 gg_id:5859 Hyphomonas oceanitis str. SCH-89 AF082
AJ007807.1 gg_id:5841 Maricaulis maris str. ACM 5106 (T) AJ0078
EU528177.1 gg_id:286163 Seasonal Dynamics Mudflat Mouth Major Kentucky Lake Reservoir Tributary sedime EU528
AJ301667.1 gg_id:26948 Maricaulis virginensis str. VC 5 AJ3016
DQ768642.1 gg_id:245137 Maricaulis virginensis MARC4M DQ768
FJ797347.1 gg_id:567500 Arenicola marina bioturbated mesocosm marine sediment clone 20 FJ7973
AY162105.1 gg_id:135262 Cultivating Sargasso Sea clone GMD21C01 AY162
NR_025323.1 gg_id:545614 Maricaulis parjimensis str. MCS 25; DSM 16078 NR_02
AJ309862.1 gg_id:40525 Oceanicaulis alexandrii str. C116-18 AJ3098
DQ644012.1 gg_id:236476 Production cyanobacterial toxin Microcystin disease corals black band diseased (BB DQ644
AY701470.1 gg_id:108547 dinoflagellate symbiont clone GCDE08_V AY701
AJ227811.1 gg_id:5847 Caulobacter sp. str. MCS33 AJ2278
GQ441337.1 gg_id:593605 and dynamic mat present vs. active nitrogen-fixing microorganisms marine microbia GQ441
FJ152756.1 gg_id:347767 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX5A_ FJ1527
DQ395426.1 gg_id:154213 deep-sea octacoral clone ctg_CGOAB48 DQ395
FJ203305.1 gg_id:307703 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2033
AB286621.1 gg_id:209532 Reducing production and domination comamonadaceae reducing oxygen supply was AB286
AF359547.1 gg_id:18991 Scrippsiella trochoidea NEPCC 15 AF359
AY942774.1 gg_id:148699 but species richness marine sponge Cymbastela concentrica clone Cc118 AY942
FJ497641.1 gg_id:542217 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4976
EU881181.1 gg_id:303607 Karst region natural restoration land soil clone KNR200711-027 EU881
AF234756.1 gg_id:18609 sludge clone S36 AF2347
GU118388.1 gg_id:538371 Threatened Corals Provide Microbial Habitats clone Gven_P10 GU118
EU196330.1 gg_id:244514 Investigation heterotrophic and autotrophic microbial perennial springs cold saline (EU196
EU245307.1 gg_id:241958 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-H5-G02 EU245
DQ376579.1 gg_id:150962 Microbial ecology removing phosphate wastewater EBPR process aerobic sequencin DQ376
AB294313.1 gg_id:205348 Microbial structure deep coal seem groundwater stream clone SWB02 AB294
AY654758.1 gg_id:105062 Oculina mucus isolate 111 AY6547
AM990862.1 gg_id:278552 Caulobacter sp. MOLA 88 AM990
FM242321.1 gg_id:337659 sequences sediment clone 83 T9d-oil FM242
AJ578477.1 gg_id:102289 Woodsholea maritima str. CM251 AJ5784
AB286463.1 gg_id:208631 Reducing production and domination comamonadaceae reducing oxygen supply was AB286
EF495229.1 gg_id:222908 Croceitomaculum antarcticum str. IMCC3195 EF4952
AJ617875.1 gg_id:110283 oxic-anoxic interphase flooded paddy soil clone D14429 AJ6178
FM242450.1 gg_id:332051 sequences sediment clone 21 T12d+oil FM242
EU665092.1 gg_id:571737 Soil are Changed Agricultural Practices agricultural soil clone Plot22-2G04 EU665
GQ500759.1 gg_id:590926 Surveys Biofilms and Sediments Mammoth Cave Karst Aquifers Bio-Sep beads conta GQ500
FJ826144.1 gg_id:514038 Succession during spring Sea filtered surface sea water decay period after diatom blo FJ8261

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FJ516864.1 gg_id:565583 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled lower s FJ5168
GU444088.1 gg_id:530619 Effects amendments on fertility and microbial structure cucumber Cucumis sativus L GU444
EU491552.1 gg_id:263491 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR4C EU491
EU236280.1 gg_id:296183 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg5aCo7 EU236
EU236341.1 gg_id:298881 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg5a2G5 EU236
EU104323.1 gg_id:309713 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
GQ441341.1 gg_id:581650 and dynamic mat present vs. active nitrogen-fixing microorganisms marine microbia GQ441
NR_026060.1 gg_id:570345 Amaricoccus tamworthensis str. Ben103; ACM 5097 NR_02
U88042.1 gg_id:5811 Amaricoccus macauensis str. Ben104 U8804
DQ130031.1 gg_id:151324 microbial biota normal forearm human skin clone GL97 DQ130
NR_029203.1 gg_id:580499 Amaricoccus veronensis str. Ben102; ACM 5098 NR_02
FJ152897.1 gg_id:336456 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX4CI FJ1528
DQ346506.1 gg_id:154801 during composting libraries compost clone 2B22 DQ346
FJ152955.1 gg_id:323451 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX4CI FJ1529
EF489005.1 gg_id:245310 Antarctobacter sp. str. TG22 EF489
EU053204.1 gg_id:246019 Catellibacterium terrae str. QMT-4 EU053
NC_009952.1 gg_id:250733 Dinoroseobacter shibae str. DFL 12 NC_00
FJ202183.1 gg_id:343990 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2021
DQ289933.1 gg_id:167723 Microbial Carbon and Nitrogen Cycling Sediments South Atlantic Bight Permeable Sh DQ289
AF452106.1 gg_id:62868 Rhodobacter massiliensis str. Framboise AF452
DQ342310.1 gg_id:205946 Haematobacter massiliensis str. H2443 DQ342
AJ534225.1 gg_id:75300 Jannaschia helgolandensis HEL-43 AJ5342
AM990650.1 gg_id:278279 culturable oligotrophic ocean (South Pacific Gyre) sea water AM990
AJ748747.1 gg_id:110922 Jannaschia rubra str. 4SM3 AJ7487
AY906862.1 gg_id:112620 Jannaschia sp. str. CL-SP26 AY9068
EF202612.1 gg_id:200932 Jannaschia donghaensis str. DSW-17 EF2026
DQ643999.1 gg_id:159971 Jannaschia pohangensis DQ643
EU034641.1 gg_id:253055 Ketogulonicigenium vulgare str. VC-2-2 EU034
DQ915605.1 gg_id:320216 Ketogulonicigenium robustum str. NRRL B-12627 DQ915
AF136849.1 gg_id:135903 Ketogulonicigenium vulgare str. DSM 4025 AF1368
AY005463.1 gg_id:45273 Leisingera methylohalidivorans str. MB2 AY0054
AJ582229.1 gg_id:93357 Loktanella salsilacus str. LMG 22002 AJ5822
DQ533555.1 gg_id:163668 Loktanella tamlensis str. SSW-27 DQ533
FJ172067.1 gg_id:327793 Microbial athalassohaline Tirez lagoon (Spain) water clone MJZ2 FJ1720
AJ582227.1 gg_id:94079 Loktanella vestfoldensis str. LMG 22006 AJ5822
AY576770.1 gg_id:138842 Ruegeria sp. str. 3X/A02/236 AY5767
AF173971.1 gg_id:73505 Roseobacter sp. str. KT1117 AF1739
FJ172062.1 gg_id:354977 Microbial athalassohaline Tirez lagoon (Spain) water clone MJW2 FJ1720
EF659438.1 gg_id:246665 within Poole Harbour seawater clone PH-F10 EF6594
AM691092.1 gg_id:220404 Rhodobacteraceae clone EG3 AM691

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DQ397336.1 gg_id:153999 Thalassobius sp. str. UST050418-052 DQ397
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DQ344498.1 gg_id:145649 Loktanella koreensis str. GA2-M3 DQ344
AY167339.1 gg_id:81378 Roseobacter sp. str. ARK9990 AY1673
FJ203462.1 gg_id:307704 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2034
AJ505788.2 gg_id:61250 coastal sediment clone AJ5057
FJ889546.1 gg_id:563888 Phaeobacter sp. str. ZS356 FJ8895
AY654808.1 gg_id:103608 Oculina mucus isolate 78 AY6548
AY654838.1 gg_id:100870 Oculina mucus isolate 40 AY6548
AF170751.1 gg_id:5658 Roseobacter sp. str. QSSC9-8 AF1707
DQ514326.1 gg_id:161345 Maribius bermudensis str. B5-6 DQ514
AY162102.1 gg_id:135261 Cultivating Sargasso Sea clone GMD21D01 AY1621
X78313.1 gg_id:5670 Ruegeria algicola str. ATCC 51441ML4 X78313
AB491842.1 gg_id:557828 affiliation (Litopenaeus vannamei) Banyuwangi East Java microbial flocs floating brac AB4918
AJ633967.1 gg_id:114827 host's accessory nidamental gland clone T65ANG1 AJ6339
FJ716854.1 gg_id:541703 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United Kingc FJ7168
AJ295988.1 gg_id:26995 Ruegeria aff. gelatinovorans str. ch611 AJ2959
EU592330.1 gg_id:539299 Seasonal changes Salton Sea hypersaline water clone SSW55N EU5923
AY258078.1 gg_id:135688 Ruegeria sp. str. DG882 AY2580
AY344411.1 gg_id:91079 Lake Kauhako surface clone K2-S-32 AY3444
GU324080.1 gg_id:556772 Marivita sp. str. CB 1101 GU324
AY600300.1 gg_id:103542 Loktanella hongkongensis str. UST950701-009P AY6003
FN431787.1 gg_id:544367 Loktanella hongkongensis str. AMV3 AMV18 FN4317
AM983542.1 gg_id:537512 Loktanella pyoseonensis str. JJM85 AM983
EU143345.1 gg_id:252067 Loktanella sp. str. J20 EU1433
AM944522.1 gg_id:268667 Nautella italica str. R-28753 AM944
FJ202078.1 gg_id:310849 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2020
FJ202115.1 gg_id:343849 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2021
EU592375.1 gg_id:583983 Seasonal changes Salton Sea hypersaline water clone SSW89Au EU5923
EF092158.1 gg_id:267120 Persistent Axinella corrugata Roseobacter clone AC2_A2 EF0921
DQ675021.1 gg_id:162482 Pseudoruegeria aquimaris str. SW-255 DQ675
DQ917854.1 gg_id:219732 Microbial profiles bleached and wild-type Muricea elongata clone BME80 DQ917
FJ202604.1 gg_id:318096 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2026
FJ202395.1 gg_id:319436 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2023
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DQ915614.1 gg_id:327289 Oceanibulbus indolifex str. DSM14862 DQ915
NR_027563.1 gg_id:544041 Oceanibulbus indolifex HEL-45 str. Hel-45; DSM 14862 NR_02
EU694388.1 gg_id:293008 Oceanibulbus indolifex str. D15 EU6943

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DQ915616.1 gg_id:305517 Oceanicola granulosus str. KCTC12143 DQ915
GQ250665.1 gg_id:544324 bulk sea water column Gulf Mexico clone GOM_WB10-7 GQ250
AB166982.1 gg_id:101280 Roseobacter sp. str. NT N37 AB166
FJ230846.1 gg_id:341497 marine str. SD7 FJ2308
FJ196062.1 gg_id:310847 Octadecabacter sp. str. ZS6-9 FJ1960
AB180384.1 gg_id:109216 Octadecabacter sp. str. No. 49 AB180
AF030437.1 gg_id:5737 Methylarcula terricola str. h37; VKM B-2160 AF0304
AF030436.1 gg_id:5762 Methylarcula terricola str. h1; VKM B-2159 AF0304
DQ402044.1 gg_id:150822 Paracoccus alcaliphilus str. mku-E2 DQ402
DQ469585.1 gg_id:158059 Paracoccus alcaliphilus str. Sty2-VH DQ469
D32240.1 gg_id:5761 Paracoccus aminovorans str. JCM 7685 D3224
GQ497295.1 gg_id:552938 Paracoccus sp. str. KSL5401-001 GQ497
GU201846.1 gg_id:518236 Paracoccus aminophilus str. Zw-11 GU201
AY312056.1 gg_id:88947 Paracoccus sp. Ol18 AY312
AF376134.1 gg_id:24018 Paracoccus sp. str. T231 AF376
FM207553.1 gg_id:348847 Paracoccus denitrificans E4 FM207
FJ381982.1 gg_id:582190 Potentially pathogenic and air stem cell transplant winter hospital shower water BMT FJ3819
EF070124.1 gg_id:190880 Paracoccus sp. TRP EF0701
DQ539490.1 gg_id:169913 Paracoccus sp. SBRD2 DQ539
DQ270723.1 gg_id:142241 Pseudomonas sp. B-1015 DQ270
AF139992.1 gg_id:5771 Paracoccus marcusii str. ML-003 AF1399
AJ582185.1 gg_id:93958 soil near uranium mill tailings clone Gitt-GS-149 AJ5821
AY014175.1 gg_id:47316 Paracoccus solventivorans str. ATCC 700252 AY014
FJ950555.1 gg_id:410450 Paracoccus versutus str. c75 FJ9505
EU434343.1 gg_id:275527 Paracoccus versutus str. a266 EU4343
DQ512861.1 gg_id:159464 Paracoccus sp. str. LW36 DQ512
AY004210.1 gg_id:28386 Paracoccus versutus str. kps AY0042
EU434456.1 gg_id:276317 Paracoccus versutus str. a234 EU4344
FM872636.1 gg_id:369875 occupant source house floor dust clone FA04D10 FM872
AY014179.1 gg_id:46743 Paracoccus yeei str. G3060 AY014
GU108563.1 gg_id:545865 active chemolithoautotrophic prokaryotes sulfidic zone Black pelagic redoxcline dete GU108
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AB275604.1 gg_id:212709 Paracoccus kamogawaensis str. YSFL3 AB2756
AJ864466.1 gg_id:109475 Paracoccus pantotrophus str. JT001 AJ8644
GQ250442.1 gg_id:577514 Paracoccus sp. str. MJ17 GQ250
AB025189.1 gg_id:5770 Paracoccus sp. str. MBIC4018 HI7 AB0251
DQ923133.1 gg_id:180344 Paracoccus halotolerans str. YIM 90738 DQ923
D32238.1 gg_id:5766 Paracoccus alcaliphilus str. JCM 7364 D32238
DQ108402.1 gg_id:137632 Paracoccus sp. Tibet-S9a3 DQ108
FJ842690.1 gg_id:576262 Paracoccus sp. str. NII-0918 FJ8426

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AM697590.1 gg_id:225243 sequencing libraries indoor dust clone BF0002D101 AM697
DQ423482.1 gg_id:155546 Paracoccus sp. str. HN-182 DQ423
EU196530.1 gg_id:241681 Paracoccus sp. Ol18 EU1965
FJ670987.1 gg_id:527854 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6709
FJ675504.1 gg_id:532336 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6755
AM231051.1 gg_id:159385 Paracoccus sp. str. R-24342 AM231
AJ864463.1 gg_id:109472 Paracoccus thiocyanatus str. SST AJ8644
FJ675534.1 gg_id:519944 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6755
AY573045.1 gg_id:106521 Paracoccus sp. str. ARCTIC-P17 AY5730
AJ534231.1 gg_id:202277 Methylarcula sp. HEL-8 AJ5342
AB255110.1 gg_id:172587 corroded concrete clone SC-148 AB2551
FM874023.1 gg_id:370368 occupant source house mattress dust clone MB05C10 FM874
DQ116009.1 gg_id:130794 bovine teat canal clone D-64 DQ116
DQ337586.1 gg_id:150080 Paracoccus sp. BBTR62 DQ337
FJ774966.1 gg_id:559136 Symbionts Brown Planthopper (Homoptera: Delphacidae) symbiont Nilaparvata luger FJ7749
EU368762.1 gg_id:267622 Paracoccus sp. str. B61 Ydz-zz EU3687
DQ129626.1 gg_id:138528 microarray level air two cities Texas urban aerosol clone AKIW1050 DQ129
GQ129980.1 gg_id:370584 Genetic inventory 'JPL-SAF' during MSL mission spacecraft assembly clean room floc GQ129
DQ532252.1 gg_id:165945 Geographical Variation Microbial Burden and Spacecraft Assembly Johnson Space Ce DQ532
D32239.1 gg_id:5758 Paracoccus aminophilus str. JCM 7686 D32239
AB264130.1 gg_id:165184 Paracoccus sp. str. TDMA-10 AB2641
GQ260189.1 gg_id:588165 Paracoccus sp. str. 22-5 GQ260
D32241.1 gg_id:5779 Paracoccus kocurii str. JCM 7684 D32241
AM900779.1 gg_id:235419 Paracoccus sp. PA216 AM900
AY167832.1 gg_id:77519 Paracoccus sp. str. SAFR-029 AY1678
AB264129.1 gg_id:164471 Paracoccus sp. str. TDMA-8 AB2641
AM990798.1 gg_id:279292 Paracoccus sp. MOLA 22 AM990
EF157164.1 gg_id:237879 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF1571
FJ267566.1 gg_id:323776 Paracoccus sp. str. II_Gauze_W_10_18 FJ2675
FJ658279.1 gg_id:570280 Metalworking Fluids and Aerosols Metallurgy Plant metalworking aerosol clone Winte FJ6582
AB098589.1 gg_id:78216 Paracoccus sp. str. TUT1021 AB0985
AB185957.1 gg_id:265461 Paracoccus marinus str. KKL-A5 AB1859
DQ856507.1 gg_id:243813 intestinal microflora Chinese mitten crab (Eriocheir sinensis) clone C1I DQ856
GQ452286.1 gg_id:582083 Paracoccus sp. str. MU2A-22 GQ452
AJ580352.1 gg_id:107767 Paracoccus sp. BOG6 AJ5803
AB012914.1 gg_id:5765 Paracoccus sp. str. MBIC3345 Y4 AB0129
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AB024613.1 gg_id:5740 Paracoccus sp. str. MBIC1145 AB0246
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DQ178660.1 gg_id:138722 Salipiger bermudensis str. HTCC2601 DQ178
AY177712.2 gg_id:83687 Phaeobacter inhibens str. T5 AY177
AF365991.1 gg_id:111393 marine str. BB2AT3 AF365
AJ867253.1 gg_id:111057 Roseobacter gallaeciensis X6 AJ8672
DQ514304.1 gg_id:161029 Roseobacter sp. str. BSs20188 DQ514
AJ633943.1 gg_id:114641 host's accessory nidamental gland clone 53ANG1 AJ6339
DQ915619.1 gg_id:311058 Phaeobacter gallaeciensis str. NBRC16654 DQ915
GQ250607.1 gg_id:588747 bulk sea water column Gulf Mexico clone GOM_WB10-40 GQ250
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FJ014985.1 gg_id:334868 Phaeobacter sp. str. M8-3.2 FJ0149
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AM884343.1 gg_id:275025 Ruegeria sp. str. SCS1 AM884
GQ348550.1 gg_id:533795 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
FJ203339.1 gg_id:314415 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2033
FJ202225.1 gg_id:350338 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2022
FJ716879.1 gg_id:554782 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United Kingc FJ7168
FJ203234.1 gg_id:305694 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2032
DQ811850.1 gg_id:172275 mangrove soil clone MSB-3ax5 DQ811
EU799430.1 gg_id:308804 It's all ranking aquatic Newport Harbour RI clone 1C227029 EU799
FJ203312.1 gg_id:327709 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2033
FJ664800.1 gg_id:554245 Quantitative dynamics cells plankton-fed microbial fuel cell clone plankton_D11 FJ6648
EU642854.1 gg_id:279275 Pseudoruegeria sp. JC2496 str. JC2496; IMSNU 14150 EU642
EU246829.1 gg_id:248590 polyphasic description Pocillopora meandrina Palmyra Atoll Calcinus obscurus abdon EU246
AF384204.1 gg_id:23935 Rhodobaca bogoriensis str. CL1 AF384
EF554833.1 gg_id:224271 Rhodobaca barguzinensis str. VKM B-2406 EF5548
AF513929.1 gg_id:62508 Rhodobacter group clone LA1-B42N AF513
AJ605746.2 gg_id:148330 Rhodobacter sp. str. EL-50 AJ6057
AF454295.1 gg_id:63841 hypersaline Mono Lake clone ML812M-6 AF454
AJ132383.1 gg_id:73883 Natronohydrobacter thiooxidans str. AHO 1 sediments ly alkaline soda lakes AJ1323
DQ659236.1 gg_id:170558 Roseinatronobacter monicus str. Ros35 DQ659
AB369166.1 gg_id:243619 Microbiological Assessment Circulation Mud Fluids During First Operation Drilling De AB369
DQ432310.1 gg_id:156883 prokaryotic alkaline hypersaline lakes Wadi Natrun Egypt Lake Hamra sediment clon DQ432
AF454292.1 gg_id:63844 hypersaline Mono Lake clone ML602M-13 AF454
DQ432219.1 gg_id:156640 prokaryotic alkaline hypersaline lakes Wadi Natrun Egypt Lake UmRisha sediment cl DQ432
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CP000663.1 gg_id:219103 Rhodobacter sphaeroides str. ATCC 17025 CP000
D16427.1 gg_id:5815 Rhodobacter capsulatus str. C5 D1642
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GU370084.1 gg_id:532163 Rhodobacter sphaeroides str. P230(1) GU370
GU455302.1 gg_id:522935 Understanding short-chain fatty acids accumulation enhanced alkaline digestion: kir GU455
AF280854.1 gg_id:49391 bioreactor clone mle1-20 AF280
GU086365.1 gg_id:560393 Pseudorhodobacter sp. str. HDW-19 GU086
FN428770.1 gg_id:543416 Looking India metagenomic approach Mahananda river water clone MRM1024 FN428
EU703461.1 gg_id:377032 lakes on Eastern Tin Plateau freshwater lake altitude 4279 masl clone XZZLH1 EU703
AB355088.1 gg_id:240664 Monitoring and archaeal two heavy polluted areas lake Egypt based on sediment Ma AB355
AY212706.1 gg_id:102583 water 10 m downstream manure clone 257ds10 AY212
EU580696.1 gg_id:268039 Rhodobacter sp. str. NP25b EU580
GQ500826.1 gg_id:587231 Surveys Biofilms and Sediments Mammoth Cave Karst Aquifers Bio-Sep beads conta GQ500
DQ815272.1 gg_id:169528 zebrafish digestive tract clone aab52e09 DQ815
EU104326.1 gg_id:321409 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
GU257946.1 gg_id:566382 Rhodobacter changlensis str. HXG-D9 GU257
DQ342322.1 gg_id:214765 Rhodobacter blasticus str. ATCC 33485; KC2138 DQ342
AB077986.1 gg_id:67721 Rhodobacter gluconicum AB077
DQ676414.1 gg_id:225039 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676
AM690348.2 gg_id:225526 Rhodobacter katedanii str. JA234 AM690
EU652478.1 gg_id:277835 Catellibacterium sp. str. JPB-2.07 EU652
GU454898.1 gg_id:520640 Understanding short-chain fatty acids accumulation enhanced alkaline digestion: kir GU454
EF192900.1 gg_id:201726 sediments temperate artificial lake Rapel reservoir sediment clone Rap2_6C EF1929
AF234745.1 gg_id:35565 sludge clone A26 AF2347
FJ382199.1 gg_id:512277 Potentially pathogenic and air stem cell transplant summer hospital shower water BN FJ3821
AF035433.1 gg_id:5793 Rhodobacter apigmentum AF0354
EU464681.1 gg_id:301082 Hartmann's mountain zebra feces clone AFZEB_aaj69a12 EU4646
AY957924.1 gg_id:114895 drinking water biofilm clone B3NR69D9 AY9579
DQ337005.1 gg_id:147896 subsurface water clone EV818CFSSAHH73 DQ337
EU979475.1 gg_id:307279 Rhodobacter sp. str. CR07-8 EU9794
AB122032.1 gg_id:98295 Rhodobacter sp. str. Jip03 AB1220
D16429.1 gg_id:5800 Rhodobacter blasticus str. ATCC33485 D16429
AF368183.1 gg_id:58379 non-EBPR sludge clone SBRT155 AF3681
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AM696303.1 gg_id:217363 Rhodovulum marinum str. JA219 AM696
D16420.1 gg_id:5840 Rhodovulum sp str. MB260 D16420
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EU770422.1 gg_id:306266 Rhodovulum sulfidophilum str. sy59 EU7704

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AM748927.1 gg_id:256512 Rhodovulum kholense str. JA297 AM748
AM921780.1 gg_id:291146 Rhodovulum puniceilacus str. JA322 AM921
NR_025222.1 gg_id:555568 Albidovulum inexpectatum str. FRR-10 NR_02
AB079635.2 gg_id:62339 Rhodovulum strictum str. JCM 9221 AB079
NR_025608.1 gg_id:568416 Roseisalinus antarcticus str. EL-88; DSM 11466 NR_02
FJ593905.2 gg_id:592117 Roseivivax sp. str. sw-2 FJ5939
FJ897782.1 gg_id:543161 Roseivivax sp. str. BH87090 FJ8977
AB166989.1 gg_id:102398 Roseivivax sp. str. NT N51 AB166
GQ243710.1 gg_id:578525 consortia immobilizing nitrifying bioreactors designed penaeid and non-penaeid larv GQ243
EF512126.1 gg_id:220146 Roseobacter denitrificans str. CSQ-3 EF5121
NC_008209.1 gg_id:163757 Roseobacter denitrificans str. OCh 114 NC_00
GQ249575.1 gg_id:513025 Microbial anaerobic sediments Bay Cadiz (Spain) marine sediment: 12-14 cm layer cl GQ249
AY654757.1 gg_id:104921 Oculina mucus isolate 112 AY654
AY177714.1 gg_id:83398 Rhodobacteraceae T11 AY177
DQ446103.2 gg_id:164817 Surface Mucopolysaccharide Layer and Band Microbial Mat Black Band-Diseased Sid DQ446
AJ844283.1 gg_id:160252 Rubellimicrobium thermophilum str. E-R2A-8a AJ8442
FM874293.1 gg_id:360432 occupant source house mattress dust clone MC04B03 FM874
EU338486.1 gg_id:242423 Pararubellimicrobium aerilata str. 5715S-9 EU338
EF547368.1 gg_id:218626 Rubellimicrobium sp. str. MSL 20 EF5473
DQ532142.1 gg_id:167156 Geographical Variation Microbial Burden and Spacecraft Assembly Jet Propulsion Lal DQ532
DQ532274.1 gg_id:172438 Geographical Variation Microbial Burden and Spacecraft Assembly Johnson Space Ce DQ532
EF682991.1 gg_id:243852 Eastern Mediterranean atmosphere clone F13_3E_FL EF6829
AJ968649.1 gg_id:144280 Ruegeria atlantica str. N5III AJ9686
DQ915630.1 gg_id:330249 Silicibacter lacuscaerulensis str. DSM11314 DQ915
DQ915631.1 gg_id:306399 Silicibacter pomeroyi str. DSM15171 DQ915
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AM709700.1 gg_id:254802 symbiotic Ruegeria strains cuttlefish Sepia officinalis egg yolk clone B4SOVto5 AM709
AM709703.1 gg_id:254660 symbiotic Ruegeria strains cuttlefish Sepia officinalis shell sacs clone 2SOSC91 AM709
AM930441.1 gg_id:268418 involved Tripneustes gratilla (Echinoidea) cloning and FISH host body wall lesions cl AM930
GU118794.1 gg_id:533437 Threatened Corals Provide Microbial Habitats clone Mfra_N19 GU118
FJ203171.1 gg_id:354074 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2031
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FJ203011.1 gg_id:350278 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2030
AM709701.1 gg_id:248371 symbiotic Ruegeria strains cuttlefish Sepia officinalis embryo clone B4SOEmb23 AM709
FJ403069.1 gg_id:357398 Disease- changes healthy and diseased colonies Caribbean coral Montastrea faveolat FJ4030
AJ633996.1 gg_id:114809 host's egg clone IEGG2 AJ6339
DQ357745.1 gg_id:149148 Marine Hatchery: Balance Between Pathogenic and Potentially Probiotic Strains mo DQ357
FJ716889.1 gg_id:549008 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United King FJ7168
EF629830.1 gg_id:274739 Changes marine Mycale laxissima on transfer aqua sponge tissue EF6298

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D88526.1 gg_id:5714 Ruegeria atlantica str. IAM14463 D8852
FJ403081.1 gg_id:369403 Disease- changes healthy and diseased colonies Caribbean coral Montastrea faveolat FJ4030
EU195952.1 gg_id:251321 Roseobacter sp. str. P123 EU195
DQ917807.1 gg_id:220204 Microbial profiles bleached and wild-type Muricea elongata Oleiphilus clone BME10 DQ917
FJ203260.1 gg_id:346584 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2032
AM709702.1 gg_id:256402 symbiotic Ruegeria strains cuttlefish Sepia officinalis renal appendage clone B4SOAr AM709
FJ937894.1 gg_id:534136 Ruegeria sp. str. LS36 FJ9378
DQ917855.1 gg_id:223353 Microbial profiles bleached and wild-type Muricea elongata Ruegeria clone ME108 DQ917
GQ391987.1 gg_id:522785 Silicibacter sp. str. PaD1.36a GQ391
FJ203147.1 gg_id:315763 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2031
AM709704.1 gg_id:235710 symbiotic Ruegeria strains cuttlefish Sepia officinalis shell sacs clone B6SOSCV22_3 AM709
EU907925.1 gg_id:328931 Ruegeria sp. str. SKUK MB1010 EU907
U58356.1 gg_id:5685 Sagittula stellata U5835
AJ557871.1 gg_id:84036 Arctic deep sea Isolation common chemoorganotrophic oxygen-respiring polar current AJ5578
FJ425632.1 gg_id:362713 communnities healthy and diseased colonies Caribbean coral Montastrea faveolata S FJ4256
FJ203292.1 gg_id:321452 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2032
DQ915633.1 gg_id:351778 Sulfitobacter brevis str. DSM11443 DQ915
DQ915634.1 gg_id:345214 Sulfitobacter delicatus str. LMG20554 DQ915
AM990658.1 gg_id:278280 Sulfitobacter sp. MOLA 381 AM990
DQ357749.1 gg_id:146001 Sulfitobacter sp. G5-5b DQ357
DQ357751.1 gg_id:150184 Sulfitobacter sp. Cl-H6 DQ357
AF007254.2 gg_id:5638 Sulfitobacter sp. str. EE-36 AF007
NZ_AALV01000004.1 gg_id:149068 Sulfitobacter sp. str. EE-36 NZ_AA
AY794211.1 gg_id:109147 soil clone F457 AY7942
GU061076.1 gg_id:592471 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061
GU061073.1 gg_id:590614 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061
GU061065.1 gg_id:544093 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061
EU823295.1 gg_id:310853 Sulfitobacter sp. str. P64 EU823
AF007260.2 gg_id:5681 Sulfitobacter sp. str. GAI-37 AF007
AF173972.1 gg_id:71580 Roseobacter sp. str. KT0917 AF173
GU061114.1 gg_id:579705 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone YAF- GU061
GU061035.1 gg_id:537289 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone Mix GU061
AJ534237.1 gg_id:75228 Sulfitobacter sp. LM-16 AJ5342
GU061104.1 gg_id:592909 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061
GU061041.1 gg_id:571091 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone Mix- GU061
AY162107.1 gg_id:135250 Cultivating Sargasso Sea clone GMD16F09 AY162
GU061096.1 gg_id:572670 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061
AY771778.1 gg_id:144788 Arctic Sulfitobacter brevis clone SE96 AY7717
EU919845.1 gg_id:315930 Kongsfjorden western Spitsbergen Svalbard ocean water clone s81 EU919
GU061088.1 gg_id:593911 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061

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AJ534236.1 gg_id:76438 Sulfitobacter sp. LM-11 AJ5342
DQ659417.1 gg_id:166927 Roseobacter sp. str. SO3 DQ659
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AY690671.1 gg_id:107730 Sulfitobacter sp. str. GC02 AY6906
AB121782.1 gg_id:104397 Jannaschia cystaugens str. CFPB-A9 AB1217
DQ915638.1 gg_id:316408 Thalassobacter stenotrophicus str. CECT5294 DQ915
AB121783.1 gg_id:104544 Jannaschia cystaugens str. CFPB-A5 AB1217
AJ631302.1 gg_id:109582 Thalassobacter stenotrophicus str. CECT 5294 = 5SM22 AJ6313
DQ915639.1 gg_id:315323 Thalassobius gelatinovorus str. NBRC15761 DQ915
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FJ202550.1 gg_id:341164 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2025
AJ968650.1 gg_id:143564 Roseovarius pelophilus str. G5II AJ9686
AF114484.2 gg_id:5704 CVSP isolate str. CV919-312 AF1144
EU236364.1 gg_id:297563 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg91D8 EU2363
FJ203176.1 gg_id:337783 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2031
FJ203129.1 gg_id:341596 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2031
DQ376573.1 gg_id:152175 Microbial ecology removing phosphate wastewater EBPR process aerobic sequencin DQ376
EF540436.1 gg_id:220130 oil-shale semi-coke solid waste clone PK_V EF5404
DQ640665.1 gg_id:233567 full-scale EBPR sludge clone Skagenf58 DQ640
EU979473.1 gg_id:344892 Rhodobacter sp. str. CR07-5 EU9794
AM403212.1 gg_id:163121 Nitrate-reducing and Denitrifying Marine Aqua Biofilter marine aqua AM403
EF205262.1 gg_id:188150 Paracoccus sp. 'CJSPY1 (P-I)' str. CJSPY1 (P-I) EF2052
FJ872532.2 gg_id:593437 Rhodobacter sp. str. ZH15 FJ8725
AY656719.1 gg_id:99632 Thioclava pacifica str. TL2 AY6567
AM745438.1 gg_id:243786 Rhodobacter maris str. JA276 AM745
AY914074.1 gg_id:114095 Rhodobacter sp. R-8 AY9140
GQ355283.1 gg_id:593578 Thioclava sp. str. GT-5 GQ355
FJ542826.1 gg_id:530503 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone A FJ5428
AY172658.1 gg_id:77954 Rhodobacter sp. str. B-7005 AY1726
AM408117.1 gg_id:186127 Rhodobacter vinaykumarii str. JA123 AM408
EF061947.1 gg_id:193660 Xiamen China mangrove sediment clone XME33 EF0619
EU491911.1 gg_id:264288 Abundance and microbial life ocean crust deep seawater East Pacific Rise clone EPR4 EU4919
FJ202842.1 gg_id:306401 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2028
FJ497541.1 gg_id:586757 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4975
GU220746.1 gg_id:591725 Microbial mineralization and involved microbial Basin field temperature hydrotherm GU220
EU287008.1 gg_id:239638 Pacific arctic surface sediment clone B78-44 EU2870
FJ905620.1 gg_id:568278 s Tonga-Kermadec iron oxide chimney-like structure on Volcano 19 Tonga Arc clone V FJ9056
AF468367.2 gg_id:80906 Arctic sea ice ARK10115 AF4683
EF573178.1 gg_id:231130 Coco's Island site 23 marine clone S23_1277 EF5731
FJ905703.1 gg_id:552641 s Tonga-Kermadec iron oxide sediments Volcano 1 Tonga Arc clone V1F23b FJ9057

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DQ270648.1 gg_id:141449 hydrothermal carbonate chimney clone LC1133B-90 DQ270
FJ497658.1 gg_id:584890 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4976
AB476190.1 gg_id:558885 Compositional and functional transition epibiotic microbial hydrothermal vent galath AB476
FJ497592.1 gg_id:556056 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4975
AY922245.1 gg_id:113571 whalefall clone 131839 AY922
FJ712530.1 gg_id:528280 Prokaryotic structure and sediments active submarine mud volcano (Kazan MV Sea) k FJ7125
FJ997596.1 gg_id:417573 Jannaschia sp. str. KYW137 FJ9975
GU061174.1 gg_id:542278 Evaluation culturing methods cultivate previously Yel Sea intertidal beach surface se GU061
EU652595.1 gg_id:278420 Seasonal variation microbial Yel Sea sediment clone D8S-60 EU652
EU919756.1 gg_id:336543 Kongsfjorden western Spitsbergen Svalbard ocean water clone b9 EU919
FJ497563.1 gg_id:586645 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4975
AM997674.1 gg_id:545505 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
GU061093.1 gg_id:584111 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061
AY794100.1 gg_id:109050 soil clone F2C01 AY794
GU119343.1 gg_id:536065 Threatened Corals Provide Microbial Habitats reef water clone Reef_C17 GU119
FJ202553.1 gg_id:341986 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2025
DQ103611.1 gg_id:135996 hypersaline endoevaporitic microbial mat clone E6bF01 DQ103
AM745160.1 gg_id:296724 Impact oil er hydrocarbons on microbial and activity Gulf Mexico cold seep marine s AM745
EU143368.1 gg_id:258074 Thalassobius sp. str. J86 EU143
AY573044.1 gg_id:106356 Roseobacter sp. str. ARCTIC-P4 AY573
AF468276.1 gg_id:80486 Arctic pack ice; northern Fram Strait; 80 31.1 N; 01 deg 59.7 min E clone ARKIA-77 AF468
AB355073.1 gg_id:242934 Monitoring and archaeal two heavy polluted areas lake Egypt based on sediment Ma AB355
DQ167248.1 gg_id:138930 Pseudoroseobacter euryhalinus str. KOPRI 13595 DQ167
FJ196006.1 gg_id:351622 Rhodobacter sp. str. ZS2-28 FJ1960
GQ349620.1 gg_id:539973 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ349
FJ203199.1 gg_id:342707 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2031
DQ395756.1 gg_id:154523 deep-sea octacoral clone ctg_CGOF276 DQ395
FJ153021.1 gg_id:308558 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX4CI FJ1530
FJ230833.2 gg_id:310112 Citricella sp. ADB str. AD8 FJ2308
FJ152760.1 gg_id:316923 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX5A_ FJ1527
FJ516861.1 gg_id:562114 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled lower s FJ5168
AJ534233.1 gg_id:75623 Staleya sp. LM-7 AJ5342
AJ534210.1 gg_id:76632 Sulfitobacter sp. DFL-10 AJ5342
AF473921.1 gg_id:41652 template extracted Diploria strigosa with BBD clone 128-27L proteobacterium AF4739
AF254109.1 gg_id:55371 hydrothermal vent strain str. TB66 AF2541
AY862771.2 gg_id:160768 prokaryotic genetic Salar de Atacama Northern Chile athalassohaline lake water clon AY8627
AJ534232.1 gg_id:75166 Staleya sp. LM-4 AJ5342
EU156067.1 gg_id:257910 Jannaschia seohaensis str. SMK-146 EU156
FJ623372.1 gg_id:591092 Efficient polyhydroxyalkanoates production waste-activated submitted process seque FJ6233
EU564842.1 gg_id:264951 Rhodovulum sp. str. CC-MHSW-4 EU5648

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DQ001322.1 gg_id:115461 Pseudorhodobacter incheonensis str. KOPRI 13537 DQ001
AM911350.1 gg_id:359759 Phenotype-specific (Scleractinia) their implications coral's nutrition health and cold- AM911
DQ395796.1 gg_id:153730 deep-sea octacoral clone ctg_CGOF245 DQ395
FJ152917.1 gg_id:310932 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX4CI FJ1529
FJ203141.1 gg_id:347865 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2031
FM242431.1 gg_id:346708 sequences sediment clone 17 T12d-oil FM242
AF353234.1 gg_id:45295 marine clone Arctic96A-8 AF3532
FJ889529.1 gg_id:560186 Sulfitobacter sp. str. KS8-4 FJ8895
FJ202949.1 gg_id:340342 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2029
GU061063.1 gg_id:543382 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061
FM958454.1 gg_id:535414 situ detection (Hippocampus guttulatus) intestinal content and mucus farmed seahc FM958
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DQ915629.1 gg_id:305904 Salipiger mucosus str. LMG22090 DQ915
EU491858.1 gg_id:272287 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU4918
FJ202508.1 gg_id:341377 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2025
AJ002565.1 gg_id:5632 Marine (isolate SRF3) clone AJ0025
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AF365993.1 gg_id:29620 marine str. BBAT2 AF3659
AF098494.1 gg_id:5645 Sulfitobacter sp. str. GAI-109 AF0984
EF202614.1 gg_id:202204 Sulfitobacter donghicola str. DSW-25 EF2026
EU573104.1 gg_id:277995 and Dependent Microbial Oil Reservoir North Sea Ekofisk oil field Norwegian sector c EU5731
AY654818.1 gg_id:99671 Oculina mucus isolate 80 AY6548
FJ202330.1 gg_id:348420 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2023
AB266065.2 gg_id:170549 Citreicella sp. str. 2-2A AB2660
GU061006.1 gg_id:564192 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone SF79 GU061
AF513400.1 gg_id:69921 Rhodobacter sp. str. 1-5 AF5134
FJ203661.1 gg_id:307998 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2036
EU544713.1 gg_id:277244 free-living particle- aggregate- archaeal and Laptev Sea 0.22-1.0 um size fraction (fre EU5447
AY162098.1 gg_id:135249 Cultivating Sargasso Sea clone GMD21H11 AY1620
DQ517132.1 gg_id:159870 Cultivation-intensive study ephemeral hypereutrophic aquatic microbial Mojave Des DQ517
EU236329.1 gg_id:301814 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg5a1F11 EU2363
AY007684.1 gg_id:41631 marine isolate JP88.1 AY0076
DQ856513.1 gg_id:245016 intestinal microflora Chinese mitten crab (Eriocheir sinensis) clone C1T DQ856
AY770942.1 gg_id:108310 water injection well Dagang oilfield clone W2 AY7709
AY922214.1 gg_id:112728 whalefall clone 131693 AY9222
AF468376.1 gg_id:79810 Arctic sea ice ARK10233 AF4683
GU061139.1 gg_id:584007 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone SF33 GU061
AB491855.1 gg_id:562026 affiliation (Litopenaeus vannamei) Banyuwangi East Java microbial flocs floating brac AB4918
GU118249.1 gg_id:592387 Threatened Corals Provide Microbial Habitats clone Dstr_B07 GU118

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GQ232737.2 gg_id:548120 Phaeobacter sp. str. G4 GQ232
FJ973591.1 gg_id:451932 Periodic sulfide irruptions impact microbial structure and hypersaline lake water colu FJ9735
DQ103619.1 gg_id:133336 hypersaline endoevaporitic microbial mat clone E6aG08 DQ103
AAYC01000011.1 gg_id:215094 Roseobacter sp. str. SK209-2-6 AAYC0
FJ202158.1 gg_id:306945 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2021
DQ395522.1 gg_id:152925 deep-sea octacoral clone ctg_CGOCA28 DQ395
DQ015795.1 gg_id:140619 Antarctic lake water clone ELB16-005 DQ015
AB114421.1 gg_id:104479 Roseovarius sp. str. S6V AB114
AY682197.1 gg_id:103418 Sulfitobacter sp. str. KMM 3457 AY682
EF646118.1 gg_id:240900 Potential interactions particle- anammox and archaeal partners namibian Namibian u EF6461
DQ856529.1 gg_id:245982 intestinal microflora Chinese mitten crab (Eriocheir sinensis) clone C2J DQ856
EF123343.1 gg_id:249977 Microbial disease microbial mats on Siderastrea siderea three regions wider Caribbea EF1233
AJ968647.1 gg_id:144272 Roseobacter sp. str. N05I AJ9686
AY701455.1 gg_id:108540 dinoflagellate symbiont clone GCHU11_G AY7014
GU061043.1 gg_id:568665 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone Mix- GU061
FJ516790.1 gg_id:575915 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled biofilm FJ5167
AY697879.1 gg_id:100581 Antarctica clone F2C25 AY6978
AF235129.1 gg_id:39391 North Sea clone KTc0993 AF2351
EU464692.1 gg_id:296206 Hartmann's mountain zebra feces clone AFZEB_aaj69c08 EU4646
EF659442.1 gg_id:247277 within Poole Harbour seawater clone PH-A6 EF6594
AM930445.1 gg_id:268773 involved Tripneustes gratilla (Echinoidea) cloning and FISH host body wall lesions clo AM930
AF507828.1 gg_id:62731 Mono Lake at depth 35 m station 6 July 2000 clone ML635J-2 proteobacterium AF5078
AM697339.1 gg_id:220866 sequencing libraries indoor dust clone BF0002C036 AM697
FJ437982.1 gg_id:542595 Benthic okenone production Fayetteville New York Green Lake surface sediments 16. FJ4379
AJ633963.1 gg_id:114571 host's accessory nidamental gland clone T63ANG236 AJ6339
AJ534243.1 gg_id:76030 Sulfitobacter sp. PIC-76 AJ5342
DQ915443.3 gg_id:168588 Maritimibacter alkaliphilus str. HTCC2654 DQ915
AY862758.2 gg_id:161659 prokaryotic genetic Salar de Atacama Northern Chile athalassohaline lake water clon AY8627
CU467448.1 gg_id:313991 Prokaryotic Tunisian saltern Saltern solar multipond clone CU467
GU117994.1 gg_id:591140 Threatened Corals Provide Microbial Habitats clone Acer_I13 GU117
FJ403075.1 gg_id:356799 Disease- changes healthy and diseased colonies Caribbean coral Montastrea faveolat FJ4030
FJ202782.1 gg_id:315498 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2027
FJ716900.1 gg_id:575317 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United Kingc FJ7169
FJ202529.1 gg_id:318405 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2025
FJ202470.1 gg_id:321216 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2024
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GQ433928.1 gg_id:560840 Subseafloor ocean crust microbial observatories: (F-through Osmo Colonization Syst GQ433
GU118103.1 gg_id:532669 Threatened Corals Provide Microbial Habitats clone Apal_O05 GU118
EF123406.1 gg_id:251365 Microbial disease microbial mats on Siderastrea siderea three regions wider Caribbea EF1234
Y15011.1 gg_id:5826 Rhodovulum iodosum str. N1 Y15011

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GU061120.1 gg_id:532672 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone YAF- GU061
AJ294354.1 gg_id:46595 Roseobacter isolate 667-4 clone AJ2943
EU246237.1 gg_id:257864 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-P5-F08 EU246
DQ234221.2 gg_id:142932 determined library mangrove clone DS139 DQ234
EU236288.1 gg_id:297968 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone HgCo15 EU236
EF648144.1 gg_id:239526 dynamics during produced water treatment aerobic activated sludge clone XJ91 EF6481
DQ372842.1 gg_id:153431 Pirellula and OM43 are among dominant lineages identified Oregon coast diatom bl DQ372
EU799185.1 gg_id:320827 It's all ranking aquatic Newport Harbour RI clone 1C226756 EU799
GU061099.1 gg_id:584882 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061
GU118238.1 gg_id:538100 Threatened Corals Provide Microbial Habitats clone Dstr_A04 GU118
DQ860070.1 gg_id:165645 Symbiotic aerobic anoxygenic phototrophic colonial tunicate clone L16 DQ860
DQ659411.1 gg_id:168003 Roseobacter sp. str. B11 DQ659
FJ516820.1 gg_id:548219 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled biofilm FJ5168
FJ203293.1 gg_id:333321 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2032
DQ856523.1 gg_id:257325 intestinal microflora Chinese mitten crab (Eriocheir sinensis) clone C2L DQ856
EF646123.1 gg_id:251424 Potential interactions particle- anammox and archaeal partners namibian Namibian u EF6461
EU573108.1 gg_id:279211 and Dependent Microbial Oil Reservoir North Sea Ekofisk oil field Norwegian sector c EU573
FJ203568.1 gg_id:352275 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2035
AY926462.1 gg_id:113266 Roseobacterium hispanicus str. B-33 AY9264
FJ202177.1 gg_id:355194 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2021
FJ976449.2 gg_id:548736 Shimia sp. str. SW-6 FJ9764
AB476254.1 gg_id:564346 Compositional and functional transition epibiotic microbial hydrothermal vent galath AB476
DQ811851.1 gg_id:165140 mangrove soil clone MSB-3B6 DQ811
EU336981.1 gg_id:240704 Saemankumia lutimaris str. SMK-117 EU336
EF632664.1 gg_id:245725 Unique microbial contrasting aquatic altitude Andean Altiplano (northern Chile) sedir EF6326
AY845238.1 gg_id:110731 sponge clone 310P14 AY8452
DQ009316.1 gg_id:123474 bacterioplankton clone SPOTSAPR01_5m244 DQ009
EU195951.1 gg_id:257413 Roseobacter sp. str. P81 EU195
FJ497575.1 gg_id:590937 Microbial Ecology Fe-rich Mats and Basaltic Rock American Samoa Vailulu'u Seamoun FJ4975
FJ203659.1 gg_id:336186 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2036
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DQ357735.1 gg_id:145906 Marine Hatchery: Balance Between Pathogenic and Potentially Probiotic Strains sea DQ357
FM180527.1 gg_id:323927 Roseovarius aestuarii str. B13 FM180
EU799662.1 gg_id:310648 It's all ranking aquatic Newport Harbour RI clone 1C227285 EU799
FJ716861.1 gg_id:542437 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United King FJ7168
AM990873.1 gg_id:278099 Structure and functions coastal NW Mediterranean ecosystem sea water AM990
GQ359327.1 gg_id:556328 Oceanicola sp. str. YCWB25 GQ359

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DQ925858.1 gg_id:165917 and marine microorganisms Antarctic sea water clone ANT40B10 DQ925
FJ516768.1 gg_id:545124 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled biofilm FJ5167
AY258088.1 gg_id:135761 Roseobacter sp. str. DG942 AY258
AY845236.1 gg_id:111182 sponge clone 33P7 AY8452
AY936186.1 gg_id:113980 Rhodobacteraceae clone F2C02 AY9361
DQ432370.1 gg_id:156995 prokaryotic alkaline hypersaline lakes Wadi Natrun Egypt Lake Hamra water clone W DQ432
FJ203264.1 gg_id:306783 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2032
DQ985046.1 gg_id:174396 Roseobacter sp. str. JL985 DQ985
EF648060.1 gg_id:251297 dynamics during produced water treatment aerobic activated sludge clone HB53 EF648
AM930444.1 gg_id:272692 involved Tripneustes gratilla (Echinoidea) cloning and FISH host body wall lesions cl AM930
GU061106.1 gg_id:569713 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061
FJ202168.1 gg_id:350486 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2021
DQ223018.1 gg_id:140852 Roseobacter sp. str. UAzPsJAC-1b DQ223
FJ202477.1 gg_id:350528 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2024
EU246273.1 gg_id:257210 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-P6-B06 EU246
FJ826135.1 gg_id:534609 Succession during spring Sea filtered surface sea water decay period after diatom blo FJ8261
FJ202682.1 gg_id:306222 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2026
GQ857654.1 gg_id:584365 Rhodobacter sp. str. RC5-103 GQ857
EF016462.1 gg_id:182573 Near real-time autonomous detection marine on coastal mooring Monterey Bay Calif EF0164
EU592337.1 gg_id:535793 Seasonal changes Salton Sea hypersaline water clone SSW5Ap EU5923
EU016198.1 gg_id:252028 Antarcticicola litoralis str. IMCC3295 EU016
DQ917848.1 gg_id:221374 Microbial profiles bleached and wild-type Muricea elongata Roseobacter clone BME DQ917
FJ403109.1 gg_id:359190 Disease- changes healthy and diseased colonies Caribbean coral Montastrea faveolat FJ4031
AY349460.1 gg_id:91740 Roseobacter sp. WHOI JT-08 AY3494
AY258090.1 gg_id:135908 Roseovarius sp. str. DG944 AY258
EF549587.1 gg_id:240979 Pseudoruegeria indica str. P73 EF5495
FJ716847.1 gg_id:554517 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United King FJ7168
EF471665.1 gg_id:276906 Fast Screening and Estuarine Operon Libraries whole surface water Chesapeake Bay EF4716
EU073791.1 gg_id:252125 Microbial subsurface beds and methanogenic enrichment s coal clone SH2B-1C EU073
DQ280370.1 gg_id:141539 Paracoccus sp. str. DS2 DQ280
EU488233.1 gg_id:556227 lucinid bivalve- symbiotic system: significance geochemical habitat on symbiont and EU488
DQ146982.1 gg_id:150257 Vibrio sp. str. V261 DQ146
AY162112.1 gg_id:135282 Cultivating Sargasso Sea clone GMD13C04 AY162
GU061952.1 gg_id:577629 and function microbial mesoscale eddy perturbations South China Sea clone S-DCM- GU061
AF473932.1 gg_id:50418 template extracted Diploria strigosa with BBD clone 128-47L proteobacterium AF4739
AY167262.1 gg_id:81992 Roseobacter sp. str. ANT9276a AY1672
FJ203083.1 gg_id:323475 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2030
DQ395613.1 gg_id:152062 deep-sea octacoral clone ctg_CGOF244 DQ395
AY690720.1 gg_id:106101 Pseudomonas sp. str. MK12 AY6907

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AF253467.1 gg_id:21303 Key aromatic-ring-cleaving enzyme protocatechuate 34-dioxygenase ecologically impc AF2534
FJ716902.1 gg_id:567191 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United Kingc FJ7169
EF186075.1 gg_id:194746 Rhodobacter sp. str. DQ12-45T EF1860
AM691099.1 gg_id:221022 Rhodobacteraceae clone EG11 AM691
FJ973583.1 gg_id:427025 Periodic sulfide irruptions impact microbial structure and hypersaline lake water colu FJ9735
AB294959.1 gg_id:222231 -dependent & analyses hydrothermal system off Taketomi Island Japan microbial ma AB2949
FJ203203.1 gg_id:348315 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2032
FJ623308.1 gg_id:583673 Efficient polyhydroxyalkanoates production waste-activated submitted process seque FJ6233
GQ451279.1 gg_id:548853 Fecal Yunnan snub-nosed monkey feces clone J274 GQ451
DQ269086.1 gg_id:189161 marine macro-alga surface clone DPC189 DQ269
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FJ155038.1 gg_id:305323 seasonal fluctuation coastal Bohai Bay sequencing and RFLP soil clone CC FJ1550
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DQ535898.1 gg_id:159839 Thalassobius aestuarii str. TF-212 DQ535
FJ403106.1 gg_id:362198 Disease- changes healthy and diseased colonies Caribbean coral Montastrea faveolat FJ4031
AM930417.1 gg_id:275915 involved Tripneustes gratilla (Echinoidea) cloning and FISH host body wall lesions clc AM930
EF659446.1 gg_id:240011 within Poole Harbour seawater clone PH-B1 EF6594
FJ516806.1 gg_id:513963 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled biofilm FJ5168
FJ203288.1 gg_id:322400 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2032
GU369891.1 gg_id:515404 and mats vents Arc white filamentous microbial mat photic shal hydrothermal vent GU369
AM696698.1 gg_id:217239 Rhodovulum adriaticum str. JA245 AM696
FJ202541.1 gg_id:311333 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2025
AM888193.1 gg_id:237685 Rhodobacter sp. str. THWCSN29 AM888
EF092221.1 gg_id:262333 Persistent Axinella corrugata Roseobacter clone B255_A1h EF0922
AB473971.1 gg_id:568966 sequences microorganisms Alvaneu Bad sulfur containing freshwater spring clone AS AB4739
GQ441231.1 gg_id:577508 and dynamic mat present vs. active nitrogen-fixing microorganisms marine microbia GQ441
DQ521519.1 gg_id:184737 Antarctic lake ice cover clone ANTLV2_E06 DQ521
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EU978825.1 gg_id:591654 and metabolic potential metagenome glacier ice clone glb337c EU9788
FJ425647.1 gg_id:364960 communnities healthy and diseased colonies Caribbean coral Montastrea faveolata R FJ4256
FJ592545.1 gg_id:549666 Fumarole-supported islands biodiversity within hyperarid -elevation landscape on Pu FJ5925
DQ811852.1 gg_id:163280 mangrove soil clone MSB-3E12 DQ811
FJ202586.1 gg_id:310049 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2025
FJ943236.1 gg_id:546187 Ponticoccus sp. IMCC5002 str. IMCC 5002 FJ9432
AM990868.1 gg_id:279616 Structure and functions coastal NW Mediterranean ecosystem sea water AM990

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EU581832.1 gg_id:546740 Oceanicola sp. str. JLT1210 EU581
AF445668.1 gg_id:21491 travertine hot spring clone SM1C11 AF445
DQ836751.1 gg_id:170685 attached biomass denitrifying bioreactor clone DB-9 DQ836
DQ223020.1 gg_id:140881 Roseobacter sp. str. UAzPsK-5 DQ223
DQ153145.1 gg_id:139458 Structure Biofilms on Artificial Surfaces and Free-living Estuarine Seawater surface b DQ153
FJ516819.1 gg_id:559891 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled biofilm FJ5168
GU118190.1 gg_id:537679 Threatened Corals Provide Microbial Habitats clone Dstr_M20 GU118
DQ071102.1 gg_id:135986 marine surface water clone SJC1.11 DQ071
FJ210801.1 gg_id:354233 Roseobacter sp. str. QDHT-05 FJ2108
AM945586.1 gg_id:263307 Sulfitobacter sp. MOLA:356 AM945
AB491831.1 gg_id:572325 affiliation (Litopenaeus vannamei) Banyuwangi East Java microbial flocs floating brac AB491
AY563033.1 gg_id:100625 Rhodobacter litoralis str. Mano9 AY563
AB100002.1 gg_id:79260 inactive deep-sea hydrothermal vent chimneys clone IndB2-5 AB100
EU799262.1 gg_id:318713 It's all ranking aquatic Newport Harbour RI clone 1C226841 EU799
AY162081.1 gg_id:135247 Cultivating Sargasso Sea clone GMD37D2 AY162
DQ860059.1 gg_id:165826 Symbiotic aerobic anoxygenic phototrophic colonial tunicate clone C3 DQ860
EU246226.1 gg_id:253444 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-P5-E03 EU246
EF123390.1 gg_id:247890 Microbial disease microbial mats on Siderastrea siderea three regions wider Caribbea EF1233
AM779569.1 gg_id:254618 Oceanicola sp. str. DongtaiB-3030 AM779
DQ015815.1 gg_id:140242 Antarctic lake water clone ELB16-059 DQ015
AJ440751.1 gg_id:66953 Gram-negative MM 1 AJ4407
EU907924.1 gg_id:335974 Roseobacter sp. str. SKUK MB1009 EU907
DQ070788.1 gg_id:146111 Prokaryotic Structures Seafloor Basalts Varying Age and Juan de Fuca Ridge deep sea DQ070
AF468352.1 gg_id:80289 Arctic sea ice ARK10013 AF4683
FJ542926.1 gg_id:515949 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone B FJ5429
EF123341.1 gg_id:239882 Microbial disease microbial mats on Siderastrea siderea three regions wider Caribbea EF1233
GQ500845.1 gg_id:533002 Surveys Biofilms and Sediments Mammoth Cave Karst Aquifers Bio-Sep beads conta GQ500
EU143346.1 gg_id:240707 Loktanella sp. str. J57 EU143
DQ446109.2 gg_id:173763 Surface Mucopolysaccharide Layer and Band Microbial Mat Black Band-Diseased Sid DQ446
GU362938.1 gg_id:548305 and archaeal biogeochemical cycling genes within Qiongdongnan basin Sea: pilot stu GU362
FJ403096.1 gg_id:365829 Disease- changes healthy and diseased colonies Caribbean coral Montastrea faveolat FJ4030
FJ825922.1 gg_id:569378 Succession during spring Sea filtered surface sea water period diatom bloom Yel Sea; FJ8259
FJ516795.1 gg_id:545503 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled biofilm FJ5167
FJ203262.1 gg_id:333817 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2032
AY922228.1 gg_id:113238 whalefall clone 131735 AY9222
FJ152875.1 gg_id:307203 Changes ly soil (Mexico) flooding alkaline saline soils former lake Texcoco clone TX4CI FJ1528
GQ349657.1 gg_id:580330 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ349
AF473915.1 gg_id:46143 template extracted Diploria strigosa with BBD clone 128-19L proteobacterium AF4739
EU246804.1 gg_id:251325 polyphasic description Pocillopora meandrina Palmyra Atoll Calcinus obscurus abdon EU2468
GU061266.1 gg_id:548114 Evaluation culturing methods cultivate previously Yel Sea intertidal beach seawater GU061

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FJ203181.1 gg_id:331968 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2031
FJ716898.1 gg_id:511895 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United Kingc FJ7168
DQ103625.1 gg_id:133177 hypersaline endoevaporitic microbial mat clone E2aE09 DQ103
EU907918.1 gg_id:345550 Roseobacter sp. str. SKUK MB1003 EU907
FJ999598.1 gg_id:435644 sponge Haliclona simulans South China Sea clone B24 FJ9995
FM872825.1 gg_id:365967 occupant source house floor dust clone FB03D09 FM872
FJ664812.1 gg_id:548236 Quantitative dynamics cells plankton-fed microbial fuel cell clone plankton_H05 FJ6648
EU592363.1 gg_id:569744 Seasonal changes Salton Sea hypersaline water clone SSW81Au EU592
DQ416564.1 gg_id:155908 s1cb87 DQ416
GU117983.1 gg_id:555923 Threatened Corals Provide Microbial Habitats clone Acer_G14 GU117
FJ716866.1 gg_id:513386 L. Bioturbated Mesocosm marine sediment Cullercoats Northumberland United Kingc FJ7168
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EU050746.1 gg_id:245675 libraries sediment Kings Bay Svalbard Arctic clone ss1_B_02_95 EU050
EU799752.1 gg_id:341789 It's all ranking aquatic Newport Harbour RI clone 1C227397 EU799
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FJ543062.1 gg_id:516831 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone B FJ5430
EF632666.1 gg_id:255072 Unique microbial contrasting aquatic altitude Andean Altiplano (northern Chile) sedir EF6326
DQ153140.1 gg_id:138819 Structure Biofilms on Artificial Surfaces and Free-living Estuarine Seawater surface b DQ153
GU118199.1 gg_id:559757 Threatened Corals Provide Microbial Habitats clone Dstr_E23 GU118
DQ421597.2 gg_id:210820 Salt Marsh Prokaryotes salt marsh clone SIMO-4232 DQ421
FJ202464.1 gg_id:323413 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2024
FJ202775.1 gg_id:343203 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2027
AB355069.1 gg_id:257450 Monitoring and archaeal two heavy polluted areas lake Egypt based on sediment Ma AB355
EU835420.1 gg_id:349125 structure and quorum sensing reverse osmosis RO membrane biofilm clone 1B15 EU835
DQ421651.2 gg_id:211048 Salt Marsh Prokaryotes salt marsh clone SIMO-4286 DQ421
DQ659415.1 gg_id:166922 Roseobacter sp. str. COL2P DQ659
AY167261.1 gg_id:81559 Roseobacter sp. str. ANT9274 AY167
EU196325.1 gg_id:243714 Roseobacter sp. NP30 EU196
EF123331.1 gg_id:239119 Microbial disease microbial mats on Siderastrea siderea three regions wider Caribbea EF1233
FJ230841.1 gg_id:304753 Phaeobacter sp. str. G8 FJ2308
DQ856532.1 gg_id:244293 intestinal microflora Chinese mitten crab (Eriocheir sinensis) clone C2N DQ856
EU835468.1 gg_id:317397 structure and quorum sensing reverse osmosis RO membrane biofilm clone 3M06 EU835
DQ234217.2 gg_id:141577 determined library mangrove clone DS134 DQ234
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DQ667965.1 gg_id:170315 Orientalimare eburneum str. SW-277 DQ667
EF207559.1 gg_id:215772 mangrove system sediment EF2075
EF471656.1 gg_id:270937 Fast Screening and Estuarine Operon Libraries whole surface water Chesapeake Bay (EF4716
FJ203039.1 gg_id:307415 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata c FJ2030

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EU685337.1 gg_id:294040 Rhodobacter sp. str. CC-SAL-30	EU6853
DQ412074.1 gg_id:155494 Sulfitobacter sp. str. SPB-4	DQ412
AY654837.1 gg_id:100715 Oculina mucus isolate 39	AY6548
Y11551.1 gg_id:5675 Roseovarius tolerans str. Ekho Lake-172	Y11551
AF076898.1 gg_id:5650 d isolate western Sargasso Sea clone D	AF0768
AY967725.1 gg_id:115231 Sulfitobacter sp. IS1	AY9677
EU287912.1 gg_id:244920 Rhodovulum thermophilus str. CC-MHSW-1	EU2879
AF359530.1 gg_id:35622 Alexandrium tamarense NEPCC 407	AF3595
DQ103624.1 gg_id:132289 hypersaline endoevaporitic microbial mat clone E4bD01	DQ103
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AY639888.1 gg_id:99168 Citricella thiooxidans str. CHLG 16	AY6398
FM242220.1 gg_id:327863 sequences sediment clone 119 T0h-oil	FM242
EU700143.1 gg_id:296248 Study on Shantou Bay Marine Sediments Microorganisms sediment clone STU1	EU7001
AF502214.1 gg_id:59351 EBPR sludge lab scale clone HP1A30	AF5022
AJ633964.1 gg_id:114588 host's accessory nidamental gland clone T63ANG237	AJ6339
AF170736.1 gg_id:5781 Rhodobacter sp. str. QSSC1-20	AF1707
AY344379.1 gg_id:92016 Lake Kauhako surface clone K2-S-3	AY3443
DQ659414.1 gg_id:167392 Roseobacter sp. str. BS90	DQ659
GQ396945.1 gg_id:593441 Functional unvegetated perhumid recently-deglaciated soils do not correlate shifts s	GQ396
EU236337.1 gg_id:295747 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg5a2D1	EU2363
AF076897.1 gg_id:5649 d isolate western Sargasso Sea clone C	AF0768
FJ202870.1 gg_id:332743 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k	FJ2028
AY922243.1 gg_id:113817 whalefall clone 131821	AY9222
EU360293.1 gg_id:271908 Pyrene-Degrading Consortium Deep-Sea Sediment West Pacific and its Key Member	EU3602
EU799036.1 gg_id:305874 It's all ranking aquatic Newport Harbour RI clone 1C226589	EU7990
DQ153126.1 gg_id:138759 Structure Biofilms on Artificial Surfaces and Free-living Estuarine Seawater surface b	DQ153
GU118282.1 gg_id:587759 Threatened Corals Provide Microbial Habitats clone Dstr_I12	GU118
EU799596.1 gg_id:350463 It's all ranking aquatic Newport Harbour RI clone 1C227214	EU7995
DQ412069.1 gg_id:156195 Sulfitobacter sp. str. KBB-2	DQ412
FJ202786.1 gg_id:352366 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k	FJ2027
EU564840.1 gg_id:259482 Pseudoruegeria sp. str. CC-GIMAT-2	EU5648
EU592378.1 gg_id:590969 Seasonal changes Salton Sea hypersaline water clone SSW59Au	EU5923
AY845243.1 gg_id:110751 sponge clone 34P35	AY8452
EU799556.1 gg_id:355768 It's all ranking aquatic Newport Harbour RI clone 1C227168	EU7995
FJ202880.1 gg_id:319087 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k	FJ2028
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FJ202552.1 gg_id:331090 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k	FJ2025
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FJ872535.1 gg_id:567410 Ruegeria sp. str. ZH17	FJ8725

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FJ906694.2 gg_id:575135 Rhodobacter sp. str. JS-43 FJ9066
EU799442.1 gg_id:354617 It's all ranking aquatic Newport Harbour RI clone 1C227041 EU7994
EF471465.1 gg_id:276208 Fast Screening and Estuarine Operon Libraries whole surface water Chesapeake Bay (EF4714
FJ232451.1 gg_id:368703 Phaeobacter sp. str. NH52F FJ2324
EF659432.1 gg_id:239375 within Poole Harbour seawater clone PH-B6 EF6594
AY845237.1 gg_id:111058 sponge clone 310P5 AY8452
GQ349664.1 gg_id:579002 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ349
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EU419924.1 gg_id:273372 Roseobacter sp. str. RW37 EU4199
GU396991.1 gg_id:516703 Haematobacter sp. str. BC14248 GU396
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EU771820.1 gg_id:337345 African elephant feces clone AFEL_aai29h10 EU7718
GU369903.1 gg_id:528643 and mats vents Arc white filamentous microbial mat photic shal hydrothermal vent GU369
DQ357748.1 gg_id:149884 Marine Hatchery: Balance Between Pathogenic and Potentially Probiotic Strains Fuc DQ357
AM176846.1 gg_id:160274 Recovery mangrove sediment clone SZB5 AM176
GU061078.1 gg_id:580648 Evaluation culturing methods cultivate previously Yel Sea intertidal beach clone TFS- GU061
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AB052711.1 gg_id:41554 Acetobacter cibinongensis str. 1BM-18C AB0527
AB052710.1 gg_id:29322 Acetobacter cibinongensis str. 4H-1 AB0527
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FJ157232.1 gg_id:318111 Acetobacter indonesiensis str. KS9 FJ1572
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AB470917.3 gg_id:535203 Acetobacter orientalis KL13-3 AB4709
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AF127399.1 gg_id:4453 Acetobacter aceti SRI1321 AF1273
NR_025512.1 gg_id:561758 Acetobacter cerevisiae str. LMG 1625 NR_025
GQ240641.1 gg_id:551528 Acetobacter pasteurianus str. DSM 2324 GQ240
EF528281.1 gg_id:224755 Acetobacter pasteurianus str. CICCHLJ Q61 EF5282
EU131161.1 gg_id:241962 Acetobacter oeni str. B7 EU1311
DQ168837.1 gg_id:138760 Fermentative biohydrogen production heated anaerobic sludge clone J20 DQ168
EU126660.1 gg_id:240163 Acetobacter pasteurianus EU1266

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AB006711.1 gg_id:4493 Acidiphilium multivorum str. AIU 301 AB0067
DQ906056.1 gg_id:227409 nickel resistance genes metagenome Tinto River rhizosphere clone ERF-1C1 DQ906
AY495956.1 gg_id:101146 Acidiphilium sp. str. WJ52 AY4959
D30775.1 gg_id:4477 Acidiphilium organovorum str. ATCC 43141 D30775
Y18446.1 gg_id:4503 Acidiphilium cryptum str. Kuesel IFV Y18446
D30776.1 gg_id:4484 Acidiphilium rubrum str. ATCC 35905 D30776
AY632901.1 gg_id:101742 Acidiphilium symbioticum str. KM2 AY6329
FJ915152.1 gg_id:541391 Acidiphilium cryptum str. nju-AMD11 FJ9151
AM502929.1 gg_id:246693 Acidiphilium cryptum str. YT1 AM502
AF376026.2 gg_id:40217 Acidiphilium sp. str. NO-17 AF3760
DQ659235.1 gg_id:220234 acid mine drainage clone DBS-Clone60 DQ659
AM697382.1 gg_id:222512 sequencing libraries indoor dust clone BF0002C079 AM697
AM947652.1 gg_id:268023 Acidisoma tundrae str. WM1 AM947
GQ264418.1 gg_id:581747 Structure Simulated Level Waste Site simulated level waste site clone WW3_107 GQ264
AM749786.1 gg_id:243689 Acidicaldus sp. T163 AM749
AM749741.1 gg_id:253578 Isolation ing candidate division soils New Zealand steam-affected geothermal soil cl AM749
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DQ906080.1 gg_id:223846 nickel resistance genes metagenome Tinto River rhizosphere clone ERF-G1 DQ906
AB362219.2 gg_id:240252 Acidocella aluminiidurans str. AL46 AB3622
D30771.1 gg_id:4491 Acidocella aminolytica str. 101 D30771
EU360486.1 gg_id:271114 pilot biological ferrous iron oxidation mine water treatment plant clone W-Z-59 EU3604
X77468.1 gg_id:4470 Acidomonas methanolica str. LMG 1668 X77468
AB056321.1 gg_id:67745 Kozakia baliensis str. Yo-3 AB0563
AB025928.1 gg_id:52420 Asaia bogorensis str. 71 AB0259
AB230993.2 gg_id:145122 Asaia bogorensis str. BCC 15703 AB2309
AB292238.1 gg_id:247489 Asaia bogorensis AG60 AB2922
EF635915.1 gg_id:255006 Asaia sp. str. Gb16 EF6359
AB025930.1 gg_id:33826 Asaia bogorensis str. 87 AB0259
AB292240.1 gg_id:246483 Asaia krungthepensis AE76 AB2922
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AB292239.1 gg_id:239498 Asaia siamensis AE97 AB2922
AB230990.2 gg_id:149795 Asaia siamensis str. BCC 15670 AB2309
CP001189.1 gg_id:306076 Gluconacetobacter diazotrophicus 5 str. PAl 5; ATCC 49037 CP0011
DQ387435.1 gg_id:220341 Gluconacetobacter diazotrophicus str. 4L DQ387
AF127392.1 gg_id:4462 Gluconacetobacter diazotrophicus SRI1990 AF1273
AF127397.1 gg_id:4461 Gluconacetobacter diazotrophicus SRI1212 AF1273
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AF127413.1 gg_id:4430 Gluconacetobacter sacchari IF9645 AF1274
AF127406.1 gg_id:4437 Gluconacetobacter sacchari SRI1216 AF1274
AJ012698.1 gg_id:4467 Gluconacetobacter europaeus str. ZIM B028 V3 AJ0126
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AF192761.1 gg_id:36976 Gluconacetobacter azotocaptans str. CFN-Ca54 AF1927
AY958232.3 gg_id:124373 Gluconacetobacter azotocaptans DS1 AY9582
AB016865.1 gg_id:4420 Acetobacter sp AB0168
AB095100.1 gg_id:71559 Gluconacetobacter persimmonensis str. KJ145 AB0951
AF127401.1 gg_id:4460 Gluconacetobacter diazotrophicus SRI1206 AF1274
AB166741.1 gg_id:104836 Gluconacetobacter saccharivorans str. LMG 1584 AB1667
AB205220.1 gg_id:114307 Gluconacetobacter europaeus str. DSM 6160 AB2052
EU000272.1 gg_id:250024 Gluconacetobacter xylinus EU000
X80775.2 gg_id:4456 Gluconobacter cerinus str. ATCC 19441 IFO 3267 = LMG 1368 X80775
AB308276.1 gg_id:231954 Gluconobacter cerinus AB3082
DQ523496.1 gg_id:159924 Acetobacter syzygii str. A265 DQ523
AB308275.1 gg_id:231476 Gluconobacter oxydans str. LD51-2 AB3082
AB178430.1 gg_id:108207 Gluconobacter oxydans str. NBRC 12118 AB1784
AB128051.1 gg_id:98968 Gluconobacter thailandicus str. F142-1 AB1280
AB178392.1 gg_id:108166 Gluconobacter oxydans str. NBRC 3250 AB1783
AB308278.1 gg_id:229359 Gluconobacter frateurii str. CR16-1 AB3082
AB470921.2 gg_id:533805 Gluconobacter frateurii GB222-1 AB4709
EU096234.1 gg_id:274069 Gluconobacter sp. str. EW707 EU096
AB470922.2 gg_id:533831 Gluconobacter japonicus GB132-1 AB4709
X82290.1 gg_id:4464 Gluconobacter frateurii str. IFO 3264 LMG 1365 X82290
AB308277.1 gg_id:234315 Gluconobacter thailandicus str. ZN22-2 AB3082
AF173825.1 gg_id:4404 Antarctic clone LB3-94 AF1738
AY682679.1 gg_id:144291 Microbial biofilms corroding systems heating system clone 24 AY6826
EU012448.1 gg_id:255991 Roseococcus suduntuyensis str. SHET EU0124
AY150047.1 gg_id:75850 Roseomonas cervicalis str. ATCC 49957 AY1500
EU742165.1 gg_id:314674 Roseomonas cervicalis str. XTD510 EU7421
GQ047159.1 gg_id:446551 Topographical and Temporal Human Skin Microbiome skin buttock clone nbw1009d GQ047
AF227831.1 gg_id:26143 str. 47076 AF2278
GQ043367.1 gg_id:395886 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw96 GQ043
GQ043897.1 gg_id:458437 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw97 GQ043
GQ038481.1 gg_id:399250 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu4 GQ038
GQ045958.1 gg_id:387425 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw9 GQ045
GQ041000.1 gg_id:378028 Topographical and Temporal Human Skin Microbiome skin inguinal crease clone nbv GQ041

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GQ045672.1 gg_id:385279 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw! GQ045
DQ532240.1 gg_id:163716 Geographical Variation Microbial Burden and Spacecraft Assembly Johnson Space C DQ532
GQ042983.1 gg_id:383540 Topographical and Temporal Human Skin Microbiome skin buttock clone nbw1011g GQ042
GQ045701.1 gg_id:371940 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw! GQ045
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GQ043724.1 gg_id:434279 Topographical and Temporal Human Skin Microbiome skin plantar heel clone nbw97 GQ043
GQ039151.1 gg_id:441025 Topographical and Temporal Human Skin Microbiome skin axillary vault clone nbu41 GQ039
AM231587.1 gg_id:152603 Roseomonas aquatica str. TR53 AM231
GQ097336.1 gg_id:424616 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw453g04 GQ097
FJ542999.1 gg_id:517152 Microbiomic intestine earthworm Eisenia fetida fed ergovaline treatment gut clone B FJ5429
AJ315073.1 gg_id:42326 Spain:Carmona clone AJ3150
GQ097313.1 gg_id:459459 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw453d1 GQ097
DQ130024.1 gg_id:154116 microbial biota normal forearm human skin clone BL102 DQ130
AY922054.1 gg_id:114073 farm soil clone AKYG679 AY922
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GQ039238.1 gg_id:415678 Topographical and Temporal Human Skin Microbiome skin axillary vault clone nbu41 GQ039
AF227855.2 gg_id:24986 str. 96446 AF2278
EU104101.1 gg_id:309755 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
EU182887.1 gg_id:236056 Roseomonas sp. str. MH59 EU1828
EF019649.1 gg_id:204768 aspen rhizosphere clone Elev_16S_801 EF0196
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DQ990962.2 gg_id:197174 Microbial Rock Varnish Yungay Atacama Desert Chile rock varnish clone AYRV2-159 DQ990
DQ129635.1 gg_id:194558 microarray level air two cities Texas urban aerosol clone AKIW418 DQ129
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FM874262.1 gg_id:358386 occupant source house mattress dust clone MC03F05 FM874
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NR_028984.1 gg_id:587551 Roseomonas rosea str. 173-96; DSM 14916 NR_02
AJ488504.1 gg_id:64742 Teichococcus ludipueritiae str. 170-96 AJ4885
EF682998.1 gg_id:246217 Eastern Mediterranean atmosphere clone F14_5H_FL EF6829
AB114611.1 gg_id:102348 rice plant clone ER-17 AB114
DQ129249.1 gg_id:151603 microarray level air two cities Texas urban aerosol clone AKIW963 DQ129
AJ871428.1 gg_id:110099 Paracraurococcus ruber CP2C AJ8714
AF513983.1 gg_id:64695 Leahibacter roseus str. LCR1 AF5139
GQ097329.1 gg_id:458521 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw453f08 GQ097
AB009013.1 gg_id:4416 Paracraurococcus ruber str. NS102 AB009
AY150045.1 gg_id:75313 Roseomonas gilardii str. ATCC 49956 AY150
EU540200.1 gg_id:280239 profile human microbiota antecubital fossa (inner elbow) skin clone nbt247a02 EU540
AF170742.1 gg_id:4413 Paracraurococcus sp. str. QSSC5-3 AF1707

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AB445099.1 gg_id:315688 Swaminathania salitolerans str. NBRC 104436 AB4450
AB304087.1 gg_id:249455 Tanticharoenia sakaeratensis str. NBRC 103193 AB3040
AB303367.1 gg_id:257476 Ameyamaea chiangmaiensis str. BCC 15745 AB3033
EF519867.1 gg_id:267119 Ronia tepidophila str. TU-7 EF5198
D45202.1 gg_id:4575 Oleomonas sagaranensis str. HD-1 D45202
EF446228.1 gg_id:213763 Prokaryotic and Ecology Floating Macroscopic Filaments Extreme Acidic River Rio Tint EF4462
EF446204.1 gg_id:208105 Prokaryotic and Ecology Floating Macroscopic Filaments Extreme Acidic River Rio Tint EF4462
EF516844.1 gg_id:220042 grassland soil clone FCPN646 EF5168
L22481.1 gg_id:8030 Heliothis virescens testis endosymbiont L22481
AF465832.1 gg_id:79493 Rubritepida flocculans AF4658
AY234551.1 gg_id:112277 soil isolate Ellin5134 AY2345
GQ088427.1 gg_id:407742 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw3 GQ088
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AJ633644.1 gg_id:103476 Rhodovarius lipocyclicus str. CCUG 44693 AJ6336
AY370188.1 gg_id:104162 ground up bee tissues clone pAJ205 AY3701
DQ378212.1 gg_id:152090 oil-polluted soil clone F50_Pitesti DQ378
FM874498.1 gg_id:361359 occupant source house mattress dust clone MD04D08 FM874
AB240243.1 gg_id:144065 microbial structure rhizosphere Phragmites bulk soil reed bed reactor laboratory clo AB2402
FJ915153.1 gg_id:583138 Acidisphaera sp. str. nju-AMDS1 FJ9151
DQ129264.1 gg_id:138221 microarray level air two cities Texas urban aerosol clone AKIW397 DQ129
EF409853.1 gg_id:207022 acid mine drainage clone SLS-49-06 EF4098
EU861903.1 gg_id:306635 effects chronic fertilization on alpine tundra microbial implications carbon and cyclin EU8619
FJ595556.1 gg_id:532895 rock Black Canyon New Mexico: and Geochemistry rocks no varnish clone NV12_7c7 FJ5955
DQ378204.1 gg_id:151650 oil-polluted soil clone F42_Pitesti DQ378
AB110421.1 gg_id:91691 Saccharibacter floricola str. S-877 AB1104
AY913353.1 gg_id:112752 forest soil clone DUNssu146 (+7A) (OTU#150) AY9133
EU409119.1 gg_id:263028 Impacted Mine Drainage Dabaoshan Mining Area mine drainage sediment clone DBS EU4091
AM162403.1 gg_id:154314 and situ identification acidic sphagnum peat bog AM162
Y07647.2 gg_id:4411 Drentse grassland soil clone vii Y07647
EU409601.1 gg_id:276553 Gluconobacter sp. str. EW707 EU4096
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AM697086.1 gg_id:221365 sequencing libraries indoor dust clone BF0002B036 AM697
AY395443.1 gg_id:102882 pasture soil clone EB1124 AY3954
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EF520426.1 gg_id:230396 structure lakes: what controls ? acid-impacted lake clone ADK-SGh02-10 EF5204
FJ465991.1 gg_id:509891 and carbon monoxide-oxidizing across successional on recent Hawaiian 1959 Kilauea FJ4659
GQ059256.1 gg_id:458919 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw: GQ059
AJ292598.1 gg_id:43037 PCB-polluted soil clone WD248 AJ2925
AM162450.1 gg_id:151048 and situ identification acidic sphagnum peat bog clone B78 AM162
EU250247.1 gg_id:250255 acid mine drainage clone GXDC-33 EU250
AB381935.1 gg_id:263857 Rhodovastum atsumiense str. G2-11 AB381
GQ402643.1 gg_id:585535 Accessing microbial biodiversity and metabolic potential tropical -based approach pe GQ402
AM691109.1 gg_id:222002 Acetobacteraceae clone BF15 AM691
EU445203.1 gg_id:554674 Isolation Weight Forest Topsoil Metagenomic tropical forest topsoil clone R16 EU445
DQ378187.1 gg_id:152683 oil-polluted soil clone F23_Pitesti DQ378
DQ378196.1 gg_id:155293 oil-polluted soil clone F33_Pitesti DQ378
AM696849.1 gg_id:226512 sequencing libraries indoor dust clone BF0002A034 AM696
AJ867914.1 gg_id:141085 Fluctuations Significantly Influence Microbial Present Nival Lakes lake water clone C2 AJ8679
FJ466191.1 gg_id:572984 and carbon monoxide-oxidizing across successional on recent Hawaiian 1959 Kilauea FJ4661
EU861954.1 gg_id:319672 effects chronic fertilization on alpine tundra microbial implications carbon and cyclin EU861
EU250257.1 gg_id:253895 acid mine drainage clone GXDC-43 EU250
FJ228277.1 gg_id:315855 sulfate-reducing enriched during bioremediation experiment sediments (Brandenbur FJ2282
DQ532333.1 gg_id:171696 Geographical Variation Microbial Burden and Spacecraft Assembly Kennedy Space C DQ532
AF465654.1 gg_id:21795 geothermal soil clone YNPRH71B AF4656
GQ495410.1 gg_id:591379 terrestrial volcanic rocks Iceland Hnausahraun lava f clone Bas-7-52 GQ495
GQ402607.1 gg_id:509496 Accessing microbial biodiversity and metabolic potential tropical -based approach pe GQ402
AJ292611.1 gg_id:44239 PCB-polluted soil clone WD2108 AJ2926
GQ054463.1 gg_id:451765 Topographical and Temporal Human Skin Microbiome skin back clone nbw1089b08c GQ054
EF516502.1 gg_id:223286 grassland soil clone FCPT506 EF5165
FJ774959.1 gg_id:509511 Symbionts Brown Planthopper (Homoptera: Delphacidae) symbiont Nilaparvata luger FJ7749
DQ659205.1 gg_id:220971 acid mine drainage clone DBS-Clone30 DQ659
DQ833767.1 gg_id:172972 Geochemical Controls on Microbial Rainbow and Springs Yellowstone National Park DQ833
AY192273.1 gg_id:81029 Rocky Mountain alpine soil clone W2b-8C AY192
FM874399.1 gg_id:364038 occupant source house mattress dust clone MD01H12 FM874
AB286050.1 gg_id:229374 Asaia lannaensis str. BCC 15733 AB286
AB297501.1 gg_id:250834 Roseomonas sp. str. K-20 (=JCM 14634) AB297
GQ055435.1 gg_id:432032 Topographical and Temporal Human Skin Microbiome skin occiput clone nbw1115c GQ055
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GU256437.1 gg_id:580136 Azospirillum amazonense str. DSM 2787 GU256
DQ288688.1 gg_id:151705 Azospirillum brasilense str. MTCC4037 DQ288
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AY563022.1 gg_id:104372 Azospirillum sp. WAI-19 AY5630
AM922283.1 gg_id:248382 Azospirillum picis str. IMMIB Tar-3 AM922
AF521650.1 gg_id:58733 Azospirillum sp. str. Arm2-2 AF5216
DQ676344.1 gg_id:220178 Archaeal sediment and plankton freshwater pond suboxic freshwater-pond clone M DQ676
AB307662.1 gg_id:260920 clay wall used Japanese traditional buildings clone R63 AB3076
AJ238567.1 gg_id:4572 Azospirillum doebereinerae str. 63f AJ2385
DQ104848.1 gg_id:131955 Azospirillum brasilense str. A154 DQ104
DQ787330.1 gg_id:171362 Azospirillum lipoferum str. B2 DQ787
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AY265875.1 gg_id:105629 perchlorate-reducing ABL1 AY2658
EU264075.1 gg_id:344876 Azospirillum sp. str. CA6 EU2640
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AB208736.1 gg_id:115023 AEOC021 clone AB2087
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AY692040.1 gg_id:107382 UASB reactor clone I43 AY6920
AY660962.1 gg_id:101185 Azospirillum lipoferum str. CN1 AY6609
AF170353.1 gg_id:4586 Azospirillum sp. str. TTI AF1703
AY118224.1 gg_id:63341 Azospirillum sp. str. DA2-3-1 AY1182
DQ288686.1 gg_id:152173 Azospirillum brasilense str. MTCC4035 DQ288
AB508878.1 gg_id:573964 Azospirillum sp. str. TSH5 AB5088
AB487643.1 gg_id:594099 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB4876
AB508897.1 gg_id:562150 Azospirillum sp. str. TSH96 AB5088
FJ970032.1 gg_id:439656 Inquilinus sp. str. CCBAU 01442 FJ9700
AM162580.1 gg_id:153285 and role capsules Sepia officinalis host egg clone 3SOMBO87 AM162
GU118836.1 gg_id:572584 Threatened Corals Provide Microbial Habitats clone Mfra_A15 GU118
AJ244806.1 gg_id:25832 Sepia officinalis symbiont AJ2448
AM162572.1 gg_id:150158 and role capsules Sepia officinalis cuttlefish egg capsule clone 3iSOMBO24 AM162
GU118235.1 gg_id:537977 Threatened Corals Provide Microbial Habitats clone Dstr_L14 GU118
GU118741.1 gg_id:555264 Threatened Corals Provide Microbial Habitats clone Mfra_E20 GU118
AJ244796.1 gg_id:34409 Sepia officinalis symbiont AJ2447
GU118225.1 gg_id:587505 Threatened Corals Provide Microbial Habitats clone Dstr_G01 GU118

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GU118209.1 gg_id:561066 Threatened Corals Provide Microbial Habitats clone Dstr_P09 GU118
FJ203583.1 gg_id:329326 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2035
AM709743.1 gg_id:256953 symbiotic Ruegeria strains cuttlefish Sepia officinalis renal appendages clone B4SOA AM709
EF612768.1 gg_id:242309 Aquaspirillum peregrinum subsp. integrum str. LMG 5407 EF6127
AM901294.1 gg_id:328525 Phaeospirillum cystidoformans str. JA317 AM901
NR_027605.1 gg_id:543918 Magnetospirillum gryphiswaldense MSR-1 str. MSR-1; DSM 6361 NR_02
AJ488096.1 gg_id:66169 chlorobenzene-degrading consortium clone IIIA-9 AJ4880
D14433.1 gg_id:4535 Phaeospirillum fulvum str. NCIMB11762 D14433
AY171615.1 gg_id:78316 Dechlorospirillum sp. str. SN1 AY1716
AJ617837.1 gg_id:110792 oxic-anoxic interphase flooded paddy soil clone D14426 AJ6178
AB250624.1 gg_id:155751 Phaeospirillum sp. str. TUT3101 AB2506
AB307663.1 gg_id:276270 clay wall used Japanese traditional buildings clone B21 AB3076
Y17390.1 gg_id:4531 Magnetospirillum sp. str. MSM-4 Y17390
AB378599.1 gg_id:259049 dominate denitrification nitrate- succinate-ammended stable istope probing 13C fra AB3785
FJ230905.1 gg_id:306797 under Level Antibiotic Selection Pressures river water clone F56 FJ2309
AB487549.1 gg_id:580150 Microbial responsive denitrification-inducing conditions rice paddy soil clone TSNSR0 AB4875
AJ292593.1 gg_id:37406 PCB-polluted soil clone WD229 AJ2925
AB111108.1 gg_id:200462 Electrochemically assisted iron respiration grow previously iron-reducing clone LS-2 AB1111
DQ659219.1 gg_id:217481 acid mine drainage clone DBS-Clone44 DQ659
AM990911.1 gg_id:279352 Nisaea sp. MOLA 137 AM990
FJ202804.1 gg_id:310439 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2028
DQ665839.1 gg_id:162790 Nisaea denitrificans str. DR41_18 DQ665
AB074520.1 gg_id:66304 Aquaspirillum itersonii subsp. nipponicum str. IAM 14945 subsp. AB0745
DQ234645.1 gg_id:143736 subsurface water clone DR546BH1103001SAD24 DQ234
EU835446.1 gg_id:313151 structure and quorum sensing reverse osmosis RO membrane biofilm clone 2B21 EU8354
AF293000.1 gg_id:50973 ferromanganous micronodule clone MNG3 AF2930
NR_028855.1 gg_id:574444 Rhodocista pekingensis str. 3-p; JCM 11669 NR_028
FM177580.1 gg_id:313312 Rhodocista pekingensis str. JA405 FM177
AM072288.1 gg_id:138132 Rhodocista xerospirillum AM072
NC_011420.2 gg_id:326714 Rhodospirillum centenum SW str. SW; ATCC 51521 NC_01
CP000230.1 gg_id:143149 Rhodospirillum rubrum str. ATCC 11170 CP0002
D30777.1 gg_id:4516 Rhodospirillum photometricum str. E-11 D30777
D14431.1 gg_id:6144 Rhodothalassium salexigens str. ATCC35888 D14431
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GU447293.1 gg_id:524419 Rhodovibrio salinarum str. LYG30 GU447
DQ432379.1 gg_id:156957 prokaryotic alkaline hypersaline lakes Wadi Natrun Egypt Lake Hamra water clone W DQ432
DQ103606.1 gg_id:136158 hypersaline endoevaporitic microbial mat clone E4aB02 DQ103
DQ103605.1 gg_id:133137 hypersaline endoevaporitic microbial mat clone E4aF06 DQ103

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AY987846.1 gg_id:136616 Rhodovibrio sp. 2Mb1 AY9878
AM284389.1 gg_id:181835 Roseospira goa str. JA133 AM284
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AM691104.1 gg_id:225998 Rhodobacteraceae clone EG16 AM691
DQ672568.1 gg_id:172363 Skermanella sp. str. 5416T-32 DQ672
GU195651.1 gg_id:584833 Skermanella sp. str. Py-2-1 GU195
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AB240277.1 gg_id:143931 microbial structure rhizosphere Phragmites bulk soil reed bed reactor laboratory clo AB240
GU134927.1 gg_id:560804 Acid-tolerant microaerophilic oxdizing enhance Fe(III) accumulation Fe(II) oxidizing e GU134
GQ402757.1 gg_id:510201 Accessing microbial biodiversity and metabolic potential tropical -based approach p GQ402
FJ823892.1 gg_id:529891 Controlling accumulation fermentation inhibitors biorefinery recycle water microbial FJ8238
FJ823930.1 gg_id:525260 Controlling accumulation fermentation inhibitors biorefinery recycle water microbial FJ8239
AM162451.1 gg_id:156377 and situ identification acidic sphagnum peat bog clone B85 AM162
AB071665.1 gg_id:67722 Tistrella mobilis AB071
FN556563.1 gg_id:514434 Tistrella sp. str. JW16.1a FN556
AM259857.1 gg_id:212680 Spatial sponge- Mediterranean Tethya aurantium sponge cortex clone TAA-10-13 AM259
EF682992.1 gg_id:235586 Eastern Mediterranean atmosphere clone F14_1C_FL EF6829
AM743175.1 gg_id:252146 Azospirillum sp. str. AP-500 AM743
EF125412.1 gg_id:176117 soil naked tidal flat sediment near mangrove clone MSB-1F1 EF1254
DQ167235.1 gg_id:138716 Brucella sp. str. HJ114 DQ167
AJ966587.1 gg_id:138336 and screening metagenomic library 5274m depth Pacific nodule province deep sea se AJ9665
FJ746303.1 gg_id:568242 Microbial oligotrophic marine sediments: South Pacific Gyre ocean sediment 5306 m FJ7463
GQ350189.1 gg_id:545318 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
EU881096.1 gg_id:331647 Karst region primary forest soil clone KPF200711-101 EU881
EF032744.1 gg_id:198274 cyanobacterial mat in lava cave clone HAVOmat83 EF0327
AY598790.1 gg_id:102860 Germany: Lake Staffelsee clone SS31 AY5987
AF391979.1 gg_id:45882 thermal soil clone YNPFFP3 AF3919
EF494345.1 gg_id:225101 Nunnock River granitic landscape clone NR.1.078 EF4943
AM293856.1 gg_id:206331 Subaequorebacter tamlense str. HST3-12 AM293
GQ264472.1 gg_id:588466 Structure Simulated Level Waste Site simulated level waste site clone WW3_179 GQ264
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EF020098.1 gg_id:206127 aspen rhizosphere clone Elev_16S_1509 EF0200
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EU919770.1 gg_id:341359 Kongsfjorden western Spitsbergen Svalbard ocean water clone b31 EU9197
EU048669.1 gg_id:250376 and archaeal surface marine sediment Slope South China Sea clone MD2896-B9 EU0486
EU236336.1 gg_id:292058 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg5a2C6 EU2363
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AB193945.1 gg_id:108907 deep-sea vent clone Sc-EB05 AB1939
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FJ535047.1 gg_id:568244 Enhancement protein conversion volatile fatty acids accumulation during addition: ef FJ5350
EU491336.1 gg_id:261444 Abundance and microbial life ocean crust seafloor lavas Loi'hi Seamount Pisces Peak EU491
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DQ378220.1 gg_id:154249 oil-polluted soil clone F58_Pitesti DQ378
DQ640710.1 gg_id:232270 full-scale EBPR sludge clone Skagen52 DQ640
EF516955.1 gg_id:226012 grassland soil clone FCPN514 EF5169
AY326611.1 gg_id:136208 Amazon soil clone 178-2 AY3266
EF515225.1 gg_id:223603 Electricigen Enrichment MFC upf microbial fuel cell anode clone 21e11 EF5152
EU133518.1 gg_id:236426 tallgrass prarie soil clone FFCH17724 EU1335
GQ350707.1 gg_id:550871 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
FJ746316.1 gg_id:540101 Microbial oligotrophic marine sediments: South Pacific Gyre ocean sediment 5306 m FJ7463
EF125410.1 gg_id:182626 soil naked tidal flat sediment near mangrove clone MSB-1E8 EF1254
GQ264148.1 gg_id:552057 Structure Simulated Level Waste Site simulated level waste site clone WC3_89 GQ264
AF104474.1 gg_id:5172 Olavius loisae endosymbiont 3 AF1044
AY921823.1 gg_id:112944 farm soil clone AKYG694 AY9218
AJ347025.1 gg_id:60184 sponge clone TK03 AJ3470
AB248648.1 gg_id:147088 mesophilic anaerobic butyrate-degrading reactor clone BHB19 AB2486
EU223956.1 gg_id:250075 -RFLP microbes Qinghai-Tin rail soil clone 127 EU2239
EU834785.1 gg_id:587763 Microbiology Aerobic Phosphorus Removal Process aerobic phosphorus-removing se EU8347
AM259856.1 gg_id:211365 Spatial sponge- Mediterranean Tethya aurantium sponge cortex clone TAA-10-03 AM259
EF157247.1 gg_id:243543 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF1572
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EU133614.1 gg_id:253675 tallgrass prarie soil clone FFCH18567 EU1336
GQ350360.1 gg_id:584854 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
AB369165.1 gg_id:241639 Microbiological Assessment Circulation Mud Fluids During First Operation Drilling De AB3691
AF406524.1 gg_id:60649 complex marine community d at depth 200 m North Aegean Sea (Station 3) clone AEG AF4065
EU834769.1 gg_id:545652 methods extraction nucleic acid activated lab scale EBPR-activated sludge clone L24 EU8347
GU134929.1 gg_id:583250 Acid-tolerant microaerophilic oxdizing enhance Fe(III) accumulation Fe(II) oxidizing e GU134
DQ015800.1 gg_id:139714 Antarctic lake water clone ELB16-121 DQ015
DQ070828.1 gg_id:147136 Prokaryotic Structures Seafloor Basalts Varying Age East Pacific Rise and Ridge basal DQ070
GQ337284.1 gg_id:572538 Hydrography shapes biogeography deep Arctic Ocean water depth 996m clone NB0 GQ337
AY913404.1 gg_id:114085 forest soil clone DUNssu198 (-7B) (OTU#132) AY9134
AJ519623.1 gg_id:75071 Uranium mill tailings soil sample clone Sh765B-TzT-31 proteobacterium AJ5196

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FJ479421.1 gg_id:511480 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4794
GQ500775.1 gg_id:552221 Surveys Biofilms and Sediments Mammoth Cave Karst Aquifers Bio-Sep beads conta GQ500
DQ917851.1 gg_id:218252 Microbial profiles bleached and wild-type Muricea elongata Ochrobactrum clone BM DQ917
EU617825.1 gg_id:343695 Seasonal variation microbial Yel Sea sediment clone C13S-26 EU617
EU491409.1 gg_id:268557 Abundance and microbial life ocean crust seafloor lavas Hawai'i South Point X4 clone EU491
FM877531.1 gg_id:368599 Microbial several mining wastes Bulgaria uranium mill tailings clone BuhD-22 FM877
EU491319.1 gg_id:260202 Abundance and microbial life ocean crust seafloor lavas Loi'hi Seamount Pisces Peak EU491
EU202862.1 gg_id:317723 -DGGE and Approach Analyze Effect Agricultural Practices on Soil agricultural soil clo EU202
GQ347126.1 gg_id:553069 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
EU236340.1 gg_id:294380 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg5a2F3 EU236
DQ015803.1 gg_id:140565 Antarctic lake water clone ELB16-042 DQ015
EU373886.1 gg_id:270370 southern Cretan margin (eastern Mediterranean Sea) canyon and slope sediment clc EU373
GU118149.1 gg_id:577466 Threatened Corals Provide Microbial Habitats clone Dstr_F11 GU118
AB445108.1 gg_id:536442 Candidatus Monilibacter spp. common bulking filaments are members III Defluviicoc AB445
DQ917837.1 gg_id:226756 Microbial profiles bleached and wild-type Muricea elongata Rhodovibrio clone BME DQ917
DQ450764.1 gg_id:160882 Microbial Alpine Tundra Wet Meadow Soil: Chloroflexi Cold Water-Saturated satura DQ450
AJ567596.1 gg_id:88278 deep-sea sediment clone MBAE43 AJ5675
AJ536857.1 gg_id:78724 uranium mining waste pile soil sample clone JG30-KF-C3 proteobacterium AJ5368
GU118759.1 gg_id:580439 Threatened Corals Provide Microbial Habitats clone Mfra_O22 GU118
DQ451512.1 gg_id:156709 forest soil clone FAC73 DQ451
AY234650.1 gg_id:111452 soil isolate Ellin5299 AY234
AY913379.1 gg_id:114006 forest soil clone DUNssu172 (-7A) (OTU#131) AY913
GQ339163.1 gg_id:523098 Comprehensive Investigation on Iron Cycling Freshwater Seep ing Microscopy Cultiv GQ339
EU236410.1 gg_id:289382 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg92E10 EU236
GQ350571.1 gg_id:550448 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
FJ675120.1 gg_id:515036 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface FJ6751
EU048609.1 gg_id:240749 Microbial surface marine sediment Xisha Trough South China Sea clone MD2902-B2 EU048
DQ300606.1 gg_id:146291 microbial ocean's interior amplicon single fosmid library pool 130m depth HOT stati DQ300
FJ902587.1 gg_id:565674 phreatic limestone sinkholes northeastern Mexico biofilm 1m deep hydrothermal ver FJ9025
AY428766.1 gg_id:103846 activated sludge AY428
EF540431.1 gg_id:222179 oil-shale semi-coke solid waste clone MK41 EF5404
EU133493.1 gg_id:254436 tallgrass prarie soil clone FFCH8309 EU133
AJ011919.1 gg_id:5104 Roseospirillum parvum str. 930I AJ0119
EU617826.1 gg_id:319668 Seasonal variation microbial Yel Sea sediment clone D13S-112 EU617
DQ815232.1 gg_id:171372 zebrafish digestive tract clone aab51h10 DQ815
EF600578.1 gg_id:292508 soil clone E5-40 EF6005
EU617823.1 gg_id:303939 Seasonal variation microbial Yel Sea sediment clone D13S-34 EU617
GQ264051.1 gg_id:568681 Structure Simulated Level Waste Site simulated level waste site clone WC3_148 GQ264
FJ478931.1 gg_id:554483 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4789
EF125433.1 gg_id:190304 mangrove soil clone MSB-2D4 EF1254

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AY627380.1 gg_id:106889 marine clone AY627380 AY6273
AY913376.1 gg_id:112806 forest soil clone DUNssu169 (-7A) (OTU#134) AY9133
AF104473.1 gg_id:4539 Olavius loisae endosymbiont 2 AF1044
CU923620.1 gg_id:545476 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU923
EU386000.1 gg_id:275575 Biogeographical and microbes sediments on subseafloor sediment South China Sea c EU386
EU652588.1 gg_id:278603 Seasonal variation microbial Yel Sea sediment clone D8S-126 EU6525
EU800562.1 gg_id:307071 It's all ranking aquatic Delaware Bay NJ clone 2C228703 EU8005
EU369121.1 gg_id:304307 libraries oyster shell clone MBIOS-05 EU3691
AM935823.1 gg_id:552916 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM935
AY913278.1 gg_id:114037 forest soil clone DUNssu056 (-2C) (OTU#127) AY9132
EU245313.1 gg_id:244972 Miniprimer new lens viewing world hypersaline microbial mat clone MAT-CR-H5-H03 EU2453
GQ346810.1 gg_id:585746 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ346
EU656113.1 gg_id:282361 Thalassobaculum sp. P24 EU6561
AY654771.1 gg_id:105164 Oculina mucus isolate 38 AY6547
AM997954.1 gg_id:574275 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
FJ592569.1 gg_id:574659 Fumarole-supported islands biodiversity within hyperarid -elevation landscape on Pu FJ5925
GQ060714.1 gg_id:419408 Topographical and Temporal Human Skin Microbiome skin glabella clone nbw04e09 GQ060
GQ263876.1 gg_id:538639 Structure Simulated Level Waste Site simulated level waste site clone WC2_121 GQ263
EF516055.1 gg_id:225294 grassland soil clone FCPU555 EF5160
DQ395408.1 gg_id:154449 deep-sea octacoral clone ctg_CGOAA92 DQ395
EU589252.1 gg_id:262768 China Ningxia rice paddy field soil clone 1_A7 EU5892
EU675666.1 gg_id:300711 Magnetospirillum sp. str. QH-2 EU6756
GQ246307.1 gg_id:513629 Temporal and spatial China North Yel Sea sediments clone Sd1-26 GQ246
EF632901.1 gg_id:244241 Unique microbial contrasting aquatic altitude Andean Altiplano (northern Chile) fresh EF6329
EF018460.1 gg_id:210561 aspen rhizosphere clone Amb_16S_1086 EF0184
GQ337197.1 gg_id:581302 Hydrography shapes biogeography deep Arctic Ocean water depth 1000m clone CB GQ337
AF407727.1 gg_id:60574 Great Artesian Basin clone B79 AF4077
Z29637.1 gg_id:4555 Azospirillum species Z29637
GQ346973.1 gg_id:580534 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ346
DQ984631.1 gg_id:253590 YK oil-contaminated soil clone YK_106 DQ984
GQ348477.1 gg_id:583127 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
EU861390.1 gg_id:318694 marine magnetite-containing magnetic spirillum str. MMS-1 EU8613
GQ349430.1 gg_id:542625 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ349
DQ906721.1 gg_id:171043 and marine microorganisms Antarctic sea water 20 m clone AntCL1D10 DQ906
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AY660963.1 gg_id:101335 Azospirillum irakense str. Rsb1 AY6609
EF018935.1 gg_id:211514 aspen rhizosphere clone Amb_16S_1444 EF0189
DQ396323.1 gg_id:152292 deep-sea octacoral clone ctg_NISA029 DQ396
EU449569.1 gg_id:561462 Soil are Changed Agricultural Practices agricultural soil clone Plot4-A11 EU4495

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GQ214100.1 gg_id:566368 China loess clone P828 GQ214
FM201297.1 gg_id:557664 'Candidatus Cryptoprodotis polytropus' Rickettsia-Like Ciliated Protist Pseudomicrot FM201
GQ264291.1 gg_id:568633 Structure Simulated Level Waste Site simulated level waste site clone WW2_11 GQ264
GU118948.1 gg_id:545008 Threatened Corals Provide Microbial Habitats clone Past_M16 GU118
AM997985.1 gg_id:563558 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
EU373875.1 gg_id:272622 southern Cretan margin (eastern Mediterranean Sea) canyon and slope sediment clc EU3738
EU133314.1 gg_id:248945 tallgrass prarie soil clone FFCH1031 EU1333
AM934985.1 gg_id:542300 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM934
EU008565.1 gg_id:236520 Nisaea salexigens str. CZ41_10a EU0085
EU133267.1 gg_id:252811 tallgrass prarie soil clone FFCH1925 EU1332
EU133443.1 gg_id:244312 tallgrass prarie soil clone FFCH15235 EU1334
EU287138.1 gg_id:249446 Pacific arctic surface sediment clone P13-45 EU2871
GU117956.1 gg_id:542538 Threatened Corals Provide Microbial Habitats clone Acer_C02 GU117
EF516857.1 gg_id:220224 grassland soil clone FCPS408 EF5168
EF516694.1 gg_id:218380 grassland soil clone FCPO401 EF5166
FM872931.1 gg_id:367617 occupant source house floor dust clone FB04G07 FM872
GQ261313.1 gg_id:573535 Study on Sediment sediment Dongping Lake clone 1S3-20 GQ261
EU652589.1 gg_id:279075 Seasonal variation microbial Yel Sea sediment clone B8S-114 EU6525
EU586202.1 gg_id:271304 Skermanella sp. str. 10-1-101 EU5862
AF179678.1 gg_id:32841 Defluvicoccus vanus str. pilsen; Ben114 AF1796
EU805031.1 gg_id:326771 It's all ranking aquatic 250 miles Panama City clone 6C233013 EU8050
AM936395.1 gg_id:537674 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM936
GU118735.1 gg_id:580777 Threatened Corals Provide Microbial Habitats clone Mfra_G10 GU118
EF157262.1 gg_id:241911 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF1572
EU665131.1 gg_id:537780 Soil are Changed Agricultural Practices agricultural soil clone Plot22-2D03 EU6651
EU335156.1 gg_id:242184 Changes microbial metabolic and along hydrogeochemically variable profile saturate EU3351
FJ592573.1 gg_id:564942 Fumarole-supported islands biodiversity within hyperarid -elevation landscape on Pu FJ5925
EU799064.1 gg_id:318113 It's all ranking aquatic Newport Harbour RI clone 1C226620 EU7990
FJ674890.1 gg_id:516834 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6748
EU802123.1 gg_id:305536 It's all ranking aquatic Chesapeake Bay MD clone 3C003533 EU8021
GQ015990.1 gg_id:493677 Topographical and Temporal Human Skin Microbiome skin back clone nbw819g09c1 GQ015
FJ202874.1 gg_id:330657 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2028
GQ402572.1 gg_id:564337 Accessing microbial biodiversity and metabolic potential tropical -based approach p GQ402
GQ406147.1 gg_id:550141 Benthic submerged sinkhole ecosystems sediment clone -45-1 GQ406
AB447630.1 gg_id:583557 Microbial Structure and Dynamics Granules Expanded Granular Sludge Bed (EGSB) R AB4476
FJ666170.1 gg_id:568338 Unseen stowaways: tank commercial cargo ship Port Houston Texas ballast water clo FJ6661
AM997864.1 gg_id:571601 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
GU220729.1 gg_id:581688 Microbial mineralization and involved microbial Basin field temperature hydrotherm GU220
EF516586.1 gg_id:220363 grassland soil clone FCPS581 EF5165

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FJ203466.1 gg_id:353797 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2034
AY963489.1 gg_id:154486 structures and correlated ecological factors soils two evergreen broad-leaved forests AY9634
GU220737.1 gg_id:535156 Microbial mineralization and involved microbial Basin field temperature hydrotherm GU220
EU802633.1 gg_id:310636 It's all ranking aquatic Northeast Colon Panama clone 4C229990 EU802
GQ348385.1 gg_id:545576 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
CU925661.1 gg_id:555845 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU925
AM997811.1 gg_id:565394 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
EF600571.1 gg_id:302726 soil clone E5-25 EF6005
EU156155.1 gg_id:256677 thermal spring microbial and functional genes Coffee Pots Hot Spring clone pCOF_65 EU1561
EU276561.1 gg_id:363728 -DGGE and Approaches Analyze Effect Agricultural Practices on Soil agricultural soil c EU2765
EU735612.1 gg_id:328739 archaeal structures and pristine soils China oil contaminated soil Jidong Oilfield clone EU7356
EU799183.1 gg_id:313413 It's all ranking aquatic Newport Harbour RI clone 1C226754 EU7991
AF355036.1 gg_id:40077 marine clone Arctic95B-2 AF3550
GQ346952.1 gg_id:583540 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ346
EU881179.1 gg_id:309422 Karst region natural restoration land soil clone KNR200711-087 EU8811
AY555804.1 gg_id:102414 Bor Khlueng hot spring clone PK324 AY5558
EF589979.2 gg_id:228872 changes structure sediments Songhua River contaminated nitrobenzene -DGGE cloni EF5899
GQ349354.1 gg_id:577322 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ349
GU118617.1 gg_id:548705 Threatened Corals Provide Microbial Habitats clone Mfav_B20 GU118
EU491294.1 gg_id:275770 Abundance and microbial life ocean crust seafloor lavas Loi'hi Seamount Pisces Peak EU4912
AY395352.1 gg_id:101876 pasture soil clone EB1033 AY3953
EU048618.1 gg_id:240989 Microbial surface marine sediment Xisha Trough South China Sea clone MD2902-B11 EU0486
EU050763.1 gg_id:245677 libraries sediment Kings Bay Svalbard Arctic clone ss1_B_07_73 EU0507
L06455.1 gg_id:4597 magnetite-containing magnetic vibrio str. MV1 L06455
EU491108.1 gg_id:266612 Abundance and microbial life ocean crust seafloor lavas Loi'hi Seamount Pisces Peak EU4911
EF157245.1 gg_id:251478 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF1572
DQ917850.1 gg_id:222549 Microbial profiles bleached and wild-type Muricea elongata Rhodospirillaceae clone DQ917
EU445208.1 gg_id:559267 Isolation Weight Forest Topsoil Metagenomic tropical forest topsoil clone R21 EU4452
AJ581583.1 gg_id:93503 soil near uranium mill tailings clone KCM-B-24 AJ5815
GQ060531.1 gg_id:405595 Topographical and Temporal Human Skin Microbiome skin glabella clone nbw02h11 GQ060
AB289439.1 gg_id:249530 Iambacter cinctapus str. IAM 15419 AB2894
GU118285.1 gg_id:546121 Threatened Corals Provide Microbial Habitats clone Dstr_H24 GU118
AM176885.1 gg_id:161167 Recovery mangrove sediment clone SZB60 AM176
DQ070832.1 gg_id:149655 Prokaryotic Structures Seafloor Basalts Varying Age East Pacific Rise and Ridge basal DQ070
FJ744861.1 gg_id:552179 Transporter genes expressed coastal response dissolved organic carbon surface wate FJ7448
GQ246388.1 gg_id:583912 Temporal and spatial China North Yel Sea sediments clone Cm1-51 GQ246
Y12596.1 gg_id:4588 Drentse grassland soil Netherlands clone DA111 Y12596
GQ060463.1 gg_id:426164 Topographical and Temporal Human Skin Microbiome skin glabella clone nbw01g03 GQ060
DQ396275.1 gg_id:151653 deep-sea octacoral clone ctg_NISA128 DQ396
FJ203534.1 gg_id:340730 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2035
DQ450755.1 gg_id:161246 Microbial Alpine Tundra Wet Meadow Soil: Chloroflexi Cold Water-Saturated satura DQ450

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GU302497.1 gg_id:567827 Spatial structure and activity sedimentary microbial underlying Beggiatoa spp. mat h GU302
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DQ499294.1 gg_id:159606 acidic cave snottite biofilm clone CV43 DQ499
DQ811855.1 gg_id:165418 mangrove soil clone MSB-4H3 DQ811
AF355035.1 gg_id:33175 marine clone Arctic96B-9a AF3550
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FJ746319.1 gg_id:561304 Microbial oligotrophic marine sediments: South Pacific Gyre ocean sediment 5306 m FJ7463
AM420239.1 gg_id:242807 Noma subgingival plaque clone 603003(oral) AM420
EU799498.1 gg_id:346828 It's all ranking aquatic Newport Harbour RI clone 1C227104 EU7994
AJ567552.1 gg_id:88580 deep-sea sediment clone MBMPE38 AJ5675
GU118834.1 gg_id:578664 Threatened Corals Provide Microbial Habitats clone Mfra_G08 GU118
FJ197425.1 gg_id:341281 deep Pacific Ocean and Eastern Mediterranean Sea sediment clone ALVIN24a_3D_FL FJ1974
EU048622.1 gg_id:242336 Microbial surface marine sediment Xisha Trough South China Sea clone MD2902-B16 EU0486
GQ264138.1 gg_id:563019 Structure Simulated Level Waste Site simulated level waste site clone WC3_78 GQ264
AY907759.1 gg_id:136640 identification picoplankton contrasting waters Arabian Sea water clone A314902 AY9077
EU236317.1 gg_id:290364 Haliclona gellius: sponge and symbionts 13 m depth Pacific Ocean clone Hg1bD5 EU2363
EF157232.1 gg_id:257142 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF1572
GQ350356.1 gg_id:589836 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
EF515995.1 gg_id:222088 grassland soil clone FCPU464 EF5159
GQ348251.1 gg_id:594329 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
AM935509.1 gg_id:534332 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM935
FM253629.1 gg_id:336148 Rock gold mine unusually microbial biofilm clone H08-1 FM253
AJ633959.1 gg_id:114556 host's accessory nidamental gland clone 30ANG6 AJ6339
GQ354964.1 gg_id:560931 structure between two geochemically distinct springs Alum Rock Park spring clone 4 GQ354
DQ451508.1 gg_id:156725 forest soil clone FAC69 DQ451
EU491777.1 gg_id:266165 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU4917
EF018173.1 gg_id:205399 aspen rhizosphere clone Amb_16S_515 EF0181
EU592393.1 gg_id:548310 Seasonal changes Salton Sea hypersaline water clone SSW14N EU5923
GU302465.1 gg_id:545368 Spatial structure and activity sedimentary microbial underlying Beggiatoa spp. mat h GU302
EF597705.1 gg_id:239558 Metagenomics deep mediterranean warm bathypelagic habitat 3010m depth Ionian EF5977
EF516219.1 gg_id:217798 grassland soil clone FCPP413 EF5162
FJ746286.1 gg_id:571876 Microbial oligotrophic marine sediments: South Pacific Gyre ocean sediment 5306 m FJ7462
EU491571.1 gg_id:273715 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR40 EU4915
GQ350038.1 gg_id:551365 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
GQ264041.1 gg_id:541154 Structure Simulated Level Waste Site simulated level waste site clone WC3_135 GQ264
EF572964.1 gg_id:231481 Coco's Island site 23 marine clone S23_1063 EF5729
EU528234.1 gg_id:284226 Seasonal Dynamics Mudflat Mouth Major Kentucky Lake Reservoir Tributary sedime EU5282
DQ070813.1 gg_id:147895 Prokaryotic Structures Seafloor Basalts Varying Age and Juan de Fuca Ridge basalt gl DQ070
EU491885.1 gg_id:271221 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU4918

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GQ351141.1 gg_id:564010 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Eastern GQ351
FJ902662.1 gg_id:578073 phreatic limestone sinkholes northeastern Mexico biofilm 1m deep hydrothermal ver FJ9026
EU799997.1 gg_id:334995 It's all ranking aquatic Newport Harbour RI clone 1C227677 EU799
FJ755769.1 gg_id:509529 vertical archaeal water and Lake Taihu sediment clone LT-SB-B68 FJ7557
DQ378260.1 gg_id:155886 oil-polluted soil clone M43_Pitesti DQ378
EU801583.1 gg_id:322226 It's all ranking aquatic Chesapeake Bay MD clone 3C002913 EU801
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DQ906739.1 gg_id:172572 and marine microorganisms Antarctic sea water 20 m clone AntCL1H5 DQ906
EU133575.1 gg_id:256291 tallgrass prarie soil clone FFCH4971 EU133
EU386027.1 gg_id:273866 Biogeographical and microbes sediments on subseafloor sediment South China Sea c EU386
AF465655.1 gg_id:33613 geothermal soil clone YNPRH85B AF465
EF018202.1 gg_id:210884 aspen rhizosphere clone Amb_16S_548 EF0182
X99671.1 gg_id:4509 Rhodospira trueperi X99671
EU050757.1 gg_id:247141 libraries sediment Kings Bay Svalbard Arctic clone ss1_B_03_94 EU050
FJ624900.1 gg_id:514040 Influence lead contamination on boreal pine forest soil clone HF_C_27 FJ6249
DQ129334.1 gg_id:138266 microarray level air two cities Texas urban aerosol clone AKIW400 DQ129
AM936707.1 gg_id:544803 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM936
EU799402.1 gg_id:341487 It's all ranking aquatic Newport Harbour RI clone 1C226994 EU799
AB054178.1 gg_id:22043 marine sponge clone HOC44 AB054
FM242428.1 gg_id:309504 sequences sediment clone 38 T12d-oil FM242
FJ826483.1 gg_id:557922 Succession during spring diatom bloom Sea filtered surface sea water contrasting nor FJ8264
AM935865.1 gg_id:548884 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM935
EF573465.1 gg_id:229444 Coco's Island site 23 marine clone S23_1564 EF5734
AF143822.1 gg_id:72005 hydrocarbon-degrading consortium clone 4-Org1-14 AF1438
AM997828.1 gg_id:547681 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
EU799295.1 gg_id:343368 It's all ranking aquatic Newport Harbour RI clone 1C226876 EU799
AJ581592.1 gg_id:94143 soil near uranium mill tailings clone KCM-B-56 AJ5815
EU800342.1 gg_id:351874 It's all ranking aquatic Delaware Bay NJ clone 2C228398 EU800
EU799038.1 gg_id:352113 It's all ranking aquatic Newport Harbour RI clone 1C226591 EU799
EU287307.1 gg_id:256038 Pacific arctic surface sediment clone S26-7 EU287
GQ350021.1 gg_id:545223 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
EU804599.1 gg_id:355413 It's all ranking aquatic 250 miles Panama City clone 6C232548 EU804
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FJ203516.1 gg_id:319588 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata h FJ2035
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EU799960.1 gg_id:342027 It's all ranking aquatic Newport Harbour RI clone 1C227638 EU799
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EU804078.1 gg_id:344865 It's all ranking aquatic 250 miles Panama City clone 6C231969 EU8040
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AY033313.1 gg_id:68981 marine bacterioplankton clone MB12C06 AY0333
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AM936369.1 gg_id:545287 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM936
DQ513034.1 gg_id:160542 Microbial life fluids ridge flank crustal fluid clone FS142-43B-02 DQ513
EF575022.1 gg_id:234907 site S25 near Coco's Island marine clone S25_1366 EF5750
DQ071034.1 gg_id:131466 marine surface water clone Chl1.16 DQ071
EU799958.1 gg_id:320279 It's all ranking aquatic Newport Harbour RI clone 1C227636 EU7999
EU802957.1 gg_id:349161 It's all ranking aquatic Northeast Colon Panama clone 4C230356 EU8029
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AB054154.1 gg_id:36213 marine sponge clone HOC20 AB0541
U70679.2 gg_id:4573 Cape Hatteras picoplankton clone OM38 U70679
FM242318.1 gg_id:307417 sequences sediment clone 51 T9d-oil FM242
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AY744399.1 gg_id:108446 EBAC2C11 complete clone EBAC2C11 AY7443
EF573783.1 gg_id:233886 site S25 near Coco's Island marine clone S25_127 EF5737
AJ240913.1 gg_id:72555 5' clone CHAB-III-8 AJ2409
EU010169.1 gg_id:252561 Biogeography Singapore seawaters seawater clone B50 EU0101
FM242430.1 gg_id:330363 sequences sediment clone 64 T12d-oil FM242
FM242317.1 gg_id:329580 sequences sediment clone 2 T9d-oil FM242
EU800341.1 gg_id:336625 It's all ranking aquatic Delaware Bay NJ clone 2C228397 EU8003
GU061713.1 gg_id:555701 and function microbial mesoscale eddy perturbations South China Sea clone CEP-5m GU061
EF540391.1 gg_id:226996 oil-shale semi-coke solid waste clone MJK32_MK EF5403
EU799440.1 gg_id:311029 It's all ranking aquatic Newport Harbour RI clone 1C227039 EU7994
EF100695.1 gg_id:175841 Caenimicrobium bisanense str. K93 EF1006
AB110414.1 gg_id:87741 Blastochloris sulfoviridis str. RN1 AB1104
FJ169957.1 gg_id:346128 Anaplasma bovis str. Zhongxian FJ1699
FJ226454.1 gg_id:350108 Anaplasma marginale str. Ishigaki-2007 FJ2264
AF318945.1 gg_id:50430 Anaplasma ovis 1/11/2/480 sheep 3573/7 AF3189
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AJ633051.1 gg_id:99352 Anaplasma ovis Zhangjiachuan AJ6330
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AJ566640.2 gg_id:136326 Ixodes ricinus endosymbiont 1 (IricES1) AJ5666
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AY969011.1 gg_id:141855 Anaplasma phagocytophilum J4 AY9690
EU978823.1 gg_id:536819 and metabolic potential metagenome glacier ice clone glb335c EU9788
EU491849.1 gg_id:264506 Abundance and microbial life ocean crust seafloor lavas East Pacific Rise clone EPR39 EU4918
GU118498.1 gg_id:587318 Threatened Corals Provide Microbial Habitats clone Gven_P15 GU118
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EF667897.1 gg_id:256410 Long-term maintenance species-specific microbiota basal metazoan Hydra epithelium EF6678
AY527214.1 gg_id:112067 Anaplasma phagocytophilum str. Strong AY5272
FJ981659.1 gg_id:563266 Flea Vectors Plague clone XC1 FJ9816
AY969014.1 gg_id:143969 Anaplasma phagocytophilum J4 AY9690
FJ478621.1 gg_id:513411 species richness estimates obtained nearly and simulated pyrosequencing-generated FJ4786
AF497583.1 gg_id:82176 detected Haemaphysalis wellingtoni removed chicken Thailand clone Hw124 AF4975
AM997528.1 gg_id:562891 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
EF667921.1 gg_id:236767 Long-term maintenance species-specific microbiota basal metazoan Hydra epithelium EF6679
EU799164.1 gg_id:355833 It's all ranking aquatic Newport Harbour RI clone 1C226733 EU7991
AB196721.1 gg_id:152987 Anaplasma phagocytophilum str. SS33P-L AB1967
AY969013.1 gg_id:143770 Anaplasma phagocytophilum J4 AY9690
GU118630.1 gg_id:582550 Threatened Corals Provide Microbial Habitats clone Mfav_B15 GU118
GU118616.1 gg_id:569314 Threatened Corals Provide Microbial Habitats clone Mfav_P11 GU118
AF156784.1 gg_id:6042 Anaplasma platys str. Gzh981 AF1567
AF493952.1 gg_id:66159 tick-borne pathogen order Rickettsiales East: evidence ticks humans identical 2 Homo AF4939
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AF318021.1 gg_id:16712 Ehrlichia ruminantium str. Porto Henrique AF3180
M73227.1 gg_id:6026 Ehrlichia ewingii M7322
NC_007798.1 gg_id:153370 Neorickettsia sennetsu str. Miyayama NC_00
AF037210.1 gg_id:6005 Neorickettsia risticii str. SHSN-1 AF0372
AF206298.1 gg_id:17901 Ehrlichia sp. 'trout isolate' trout AF2062
AF037211.1 gg_id:6006 Neorickettsia risticii str. SHSN-2 AF0372

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X71837.1 gg_id:5918 Caedibacter caryophilus str. 221 X71837
AF132138.1 gg_id:39058 Acanthamoeba polyphaga HN-3 symbiont AF1321
DQ129376.1 gg_id:154693 microarray level air two cities Texas urban aerosol clone AKIW1012 DQ129
DQ537528.1 gg_id:173557 manganese oxidation soil clone JH-WH29 DQ537
EF575101.1 gg_id:233724 site S25 near Coco's Island marine clone S25_1445 EF5751
AM997318.1 gg_id:562327 and biogeography deep-sea surface sediments South Ocean Atlantic Ocean:South-A AM997
CU926702.1 gg_id:564680 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU926
FJ529991.1 gg_id:544240 Initial ammonia- and nitrite-oxidizing compromises successful nitritation counter-diffu FJ5299
EU735608.1 gg_id:305493 archaeal structures and pristine soils China oil contaminated soil Jidong Oilfield clone EU735
DQ925896.1 gg_id:252167 Microbes and Thermal Gradients Temperature Hydrothermal Chimneys hydrotherm DQ925
AY174894.1 gg_id:110755 Sinorickettsia chlamys AY1748
AB299552.1 gg_id:254738 Motility Symbiont Termite Gut Flagellate Caduceia versatilis Member 'Synergistes' G AB2995
DQ337066.1 gg_id:146295 subsurface water clone EV818SWSAP21 DQ337
AB185652.1 gg_id:106778 rumen clone F24-H08 AB185
AB192232.1 gg_id:141713 termite gut homogenate clone RsTu1-71 AB192
AB089088.1 gg_id:78348 termite gut homogenate clone Rs-D21 proteobacterium AB089
AB192283.1 gg_id:144338 termite gut homogenate clone RPK-86 AB192
FJ202085.1 gg_id:352689 and White Plague Disease-Induced Changes Caribbean Coral Montastraea faveolata k FJ2020
AB192282.1 gg_id:142916 termite gut homogenate clone RPK-49 AB192
AJ867903.1 gg_id:143746 Fluctuations Significantly Influence Microbial Present Nival Lakes lake water clone A1 AJ8679
EF409837.1 gg_id:208265 acid mine drainage clone YTW-34-06 EF4098
AY493937.1 gg_id:136282 Fingerprinting aggregates soil clone 15 AY4939
GQ008144.1 gg_id:501518 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbw GQ008
GQ385282.1 gg_id:566774 Biofilm formation on new membranes initiated and dominated Spinghomonas spp r GQ385
AF132139.1 gg_id:40401 Acanthamoeba endosymbiont UWET39 AF1321
AF069496.1 gg_id:4551 Candidatus Odyssella thessalonicensis str. L13 AF0694
EU305601.1 gg_id:244100 Short-term effect organic matter on nitrite oxidation performance structure nitrite-o EU305
DQ464145.1 gg_id:157472 acid mine drainage clone D3-44 DQ464
DQ336985.1 gg_id:148878 subsurface water clone EV818CFSSAHH30 DQ336
EU133464.1 gg_id:247958 tallgrass prarie soil clone FFCH1626 EU1334
FJ437938.1 gg_id:550142 Benthic okenone production Fayetteville New York Green Lake surface sediments 16. FJ4379
AB089089.1 gg_id:78233 termite gut homogenate clone Rs-D39 proteobacterium AB089
EF140635.1 gg_id:180865 Acanthamoeba sp. KA/E9 symbiont EF1406
GQ500786.1 gg_id:579026 Surveys Biofilms and Sediments Mammoth Cave Karst Aquifers Bio-Sep beads conta GQ500
AJ318202.1 gg_id:45099 waste-gas biofilter clone BIxi22 AJ3182
AF132137.1 gg_id:24244 Acanthamoeba endosymbiont UWC9 AF1321
AY135665.1 gg_id:106090 Southern Ocean clone ANT18/2_83 AY1356
EU804706.1 gg_id:348480 It's all ranking aquatic 250 miles Panama City clone 6C232664 EU8047
FJ155034.1 gg_id:341080 seasonal fluctuation coastal Bohai Bay sequencing and RFLP ocean water clone BJ FJ1550
EF572082.1 gg_id:232332 Coco's Island site 23 marine clone S23_181 EF5720
EF573154.1 gg_id:233774 Coco's Island site 23 marine clone S23_1253 EF5731

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EU010190.1 gg_id:247318 Biogeography Singapore seawaters seawater clone C4 EU010
EF572152.1 gg_id:230071 Coco's Island site 23 marine clone S23_251 EF5721
EU237277.1 gg_id:251339 Genes transcripts Eastern and Western Equatorial North Atlantic Ocean microbial as: EU237
EU394583.1 gg_id:352939 Coccolithophore bloom dynamics shape (NE-Atl) northern Bay Biscay pelagic particle EU394
EF573604.1 gg_id:228750 Coco's Island site 23 marine clone S23_1703 EF5736
EU804172.1 gg_id:347854 It's all ranking aquatic 250 miles Panama City clone 6C232068 EU804
GQ337086.1 gg_id:550700 Hydrography shapes biogeography deep Arctic Ocean water depth 1191m clone NB: GQ337
DQ071035.1 gg_id:130275 marine surface water clone Chl1.17 DQ071
EU268097.1 gg_id:249237 summer Dalian and Liaodong bay determined sea water clone L4163 EU268
EU804840.1 gg_id:322061 It's all ranking aquatic 250 miles Panama City clone 6C232806 EU804
EU804308.1 gg_id:348507 It's all ranking aquatic 250 miles Panama City clone 6C232210 EU804
GU119425.1 gg_id:538759 Threatened Corals Provide Microbial Habitats reef water clone Reef_H23 GU119
EU804403.1 gg_id:312309 It's all ranking aquatic 250 miles Panama City clone 6C232318 EU804
EU804919.1 gg_id:322681 It's all ranking aquatic 250 miles Panama City clone 6C232895 EU804
FJ744775.1 gg_id:553689 Transporter genes expressed coastal response dissolved organic carbon surface wate FJ7447
GU061340.1 gg_id:569849 and function microbial mesoscale eddy perturbations South China Sea clone CE1-5m GU061
GQ348144.1 gg_id:586176 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
EU800313.1 gg_id:349820 It's all ranking aquatic Delaware Bay NJ clone 2C228362 EU800
FJ826485.1 gg_id:562447 Succession during spring diatom bloom Sea filtered surface sea water contrasting nor FJ8264
EF572554.1 gg_id:229787 Coco's Island site 23 marine clone S23_653 EF5725
EU804543.1 gg_id:345570 It's all ranking aquatic 250 miles Panama City clone 6C232490 EU804
AF382134.1 gg_id:79390 bacterioplankton clone ZA3320c AF382
EU804583.1 gg_id:316863 It's all ranking aquatic 250 miles Panama City clone 6C232532 EU804
GU119354.1 gg_id:548940 Threatened Corals Provide Microbial Habitats reef water clone Reef_P04 GU119
EF572954.1 gg_id:234414 Coco's Island site 23 marine clone S23_1053 EF5729
FJ744853.1 gg_id:532890 Transporter genes expressed coastal response dissolved organic carbon surface wate FJ7448
EU799492.1 gg_id:352428 It's all ranking aquatic Newport Harbour RI clone 1C227098 EU799
EU237401.1 gg_id:245547 Genes transcripts Eastern and Western Equatorial North Atlantic Ocean microbial as: EU237
EU799198.1 gg_id:338982 It's all ranking aquatic Newport Harbour RI clone 1C226770 EU799
EU805287.1 gg_id:303698 It's all ranking aquatic 250 miles Panama City clone 6C233287 EU805
GQ350090.1 gg_id:575288 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
EU804214.1 gg_id:344952 It's all ranking aquatic 250 miles Panama City clone 6C232114 EU804
GU061805.1 gg_id:581967 and function microbial mesoscale eddy perturbations South China Sea clone CEP-DC GU061
FJ545626.1 gg_id:567003 Diveristy Seawater and Sediment Northern North Yel Sea seawater clone W2-50 FJ5456
GU061372.1 gg_id:553174 and function microbial mesoscale eddy perturbations South China Sea clone CE1-5m GU061
EU804569.1 gg_id:338938 It's all ranking aquatic 250 miles Panama City clone 6C232517 EU804
EU805276.1 gg_id:318071 It's all ranking aquatic 250 miles Panama City clone 6C233276 EU805
FJ744769.1 gg_id:550347 Transporter genes expressed coastal response dissolved organic carbon surface wate FJ7447
EU800579.1 gg_id:340109 It's all ranking aquatic Delaware Bay NJ clone 2C228725 EU800
EF572965.1 gg_id:230129 Coco's Island site 23 marine clone S23_1064 EF5729
EU804139.1 gg_id:353394 It's all ranking aquatic 250 miles Panama City clone 6C232032 EU804

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EU802744.1 gg_id:315894 It's all ranking aquatic Northeast Colon Panama clone 4C230111 EU8027
EF572652.1 gg_id:228725 Coco's Island site 23 marine clone S23_751 EF5726
EU010214.1 gg_id:249320 Biogeography Singapore seawaters seawater clone C8 EU0102
GU230220.1 gg_id:584057 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A GU230
EU805083.1 gg_id:327126 It's all ranking aquatic 250 miles Panama City clone 6C233068 EU8050
EU802898.1 gg_id:305898 It's all ranking aquatic Northeast Colon Panama clone 4C230290 EU8028
EU410957.1 gg_id:258240 Description SAR11 containing lighting region variable replacement within genome Ca EU4109
EU804893.1 gg_id:318781 It's all ranking aquatic 250 miles Panama City clone 6C232865 EU8048
EU804562.1 gg_id:340648 It's all ranking aquatic 250 miles Panama City clone 6C232509 EU8045
EU802304.1 gg_id:307321 It's all ranking aquatic Northeast Colon Panama clone 4C229615 EU8023
EU804721.1 gg_id:329320 It's all ranking aquatic 250 miles Panama City clone 6C232679 EU8047
GU119190.1 gg_id:554445 Threatened Corals Provide Microbial Habitats reef water clone Reef_G03 GU119
EU800774.1 gg_id:341025 It's all ranking aquatic Delaware Bay NJ clone 2C228971 EU8007
U75253.1 gg_id:5912 marine planktonic clone SAR407 U75253
EU804381.1 gg_id:355665 It's all ranking aquatic 250 miles Panama City clone 6C232293 EU8043
EF572413.1 gg_id:233290 Coco's Island site 23 marine clone S23_512 EF5724
EU804710.1 gg_id:317021 It's all ranking aquatic 250 miles Panama City clone 6C232668 EU8047
EF572687.1 gg_id:229583 Coco's Island site 23 marine clone S23_786 EF5726
EU804612.1 gg_id:350897 It's all ranking aquatic 250 miles Panama City clone 6C232561 EU8046
GQ350315.1 gg_id:592999 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
EU802940.1 gg_id:355076 It's all ranking aquatic Northeast Colon Panama clone 4C230336 EU8029
EF573232.1 gg_id:231199 Coco's Island site 23 marine clone S23_1331 EF5732
EU804547.1 gg_id:326831 It's all ranking aquatic 250 miles Panama City clone 6C232494 EU8045
FJ826220.1 gg_id:554175 Succession during spring diatom bloom Sea filtered surface sea water contrasting nor FJ8262
GQ348078.1 gg_id:593145 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
EU800420.1 gg_id:339916 It's all ranking aquatic Delaware Bay NJ clone 2C228516 EU8004
GQ346823.1 gg_id:587245 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ346
EU804281.1 gg_id:328071 It's all ranking aquatic 250 miles Panama City clone 6C232183 EU8042
GQ347719.1 gg_id:576263 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
EF571967.1 gg_id:231214 Coco's Island site 23 marine clone S23_66 EF5719
EU801166.1 gg_id:335800 It's all ranking aquatic Delaware Bay NJ clone 2C229523 EU8011
EF573156.1 gg_id:230611 Coco's Island site 23 marine clone S23_1255 EF5731
GQ350674.1 gg_id:536841 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
FJ826114.1 gg_id:565466 Succession during spring Sea filtered surface sea water decay period after diatom blo FJ8261
GQ346738.1 gg_id:538218 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ346
EF573022.1 gg_id:232263 Coco's Island site 23 marine clone S23_1121 EF5730
EF572419.1 gg_id:228811 Coco's Island site 23 marine clone S23_518 EF5724
EF571934.1 gg_id:230704 Coco's Island site 23 marine clone S23_33 EF5719
FJ744970.1 gg_id:536126 Transporter genes expressed coastal response dissolved organic carbon surface wate FJ7449
EF572311.1 gg_id:230398 Coco's Island site 23 marine clone S23_410 EF5723

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EU802469.1 gg_id:338687 It's all ranking aquatic Northeast Colon Panama clone 4C229801 EU802
GQ350531.1 gg_id:538597 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
DQ071141.1 gg_id:136072 marine surface water clone Surf1.16 DQ071
EU800481.1 gg_id:326350 It's all ranking aquatic Delaware Bay NJ clone 2C228595 EU800
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EU805349.1 gg_id:346932 It's all ranking aquatic 250 miles Panama City clone 6C233353 EU805
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EU800477.1 gg_id:331844 It's all ranking aquatic Delaware Bay NJ clone 2C228590 EU800
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EU802582.1 gg_id:316095 It's all ranking aquatic Northeast Colon Panama clone 4C229931 EU802
EU800272.1 gg_id:327151 It's all ranking aquatic Delaware Bay NJ clone 2C228312 EU800
GQ350462.1 gg_id:562464 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
DQ655710.1 gg_id:161812 ocean) brackish zone shal coastal lagoon (South Atlantic Ocean) clone 8 DQ655
DQ071057.1 gg_id:136068 marine surface water clone Chl1.44 DQ071
EF573476.1 gg_id:228126 Coco's Island site 23 marine clone S23_1575 EF5734
EF572494.1 gg_id:232394 Coco's Island site 23 marine clone S23_593 EF5724
EU805294.1 gg_id:309191 It's all ranking aquatic 250 miles Panama City clone 6C233294 EU805
EU801916.1 gg_id:339935 It's all ranking aquatic Chesapeake Bay MD clone 3C003295 EU801
EU805290.1 gg_id:336652 It's all ranking aquatic 250 miles Panama City clone 6C233290 EU805
EU592342.1 gg_id:563209 Seasonal changes Salton Sea hypersaline water clone SSW26Ap EU5923
EF574158.1 gg_id:231586 site S25 near Coco's Island marine clone S25_502 EF5741
EF572576.1 gg_id:234584 Coco's Island site 23 marine clone S23_675 EF5725
GQ348209.1 gg_id:568564 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
EU804711.1 gg_id:311847 It's all ranking aquatic 250 miles Panama City clone 6C232669 EU804
EU805387.1 gg_id:316104 It's all ranking aquatic 250 miles Panama City clone 6C233396 EU805
EU805281.1 gg_id:329803 It's all ranking aquatic 250 miles Panama City clone 6C233281 EU805
GU061691.1 gg_id:586444 and function microbial mesoscale eddy perturbations South China Sea clone CEP-5m GU061
GU061692.1 gg_id:591579 and function microbial mesoscale eddy perturbations South China Sea clone CEP-5m GU061
EU005773.1 gg_id:276591 Major activity inside and outside natural iron-fertilized phytoplankton bloom 3 micro EU005
EU800693.1 gg_id:345826 It's all ranking aquatic Delaware Bay NJ clone 2C228861 EU800
DQ071050.1 gg_id:132375 marine surface water clone Chl1.38 DQ071
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EU268085.1 gg_id:240606 summer Dalian and Liaodong bay determined sea water clone L2104 EU268
EF573262.1 gg_id:231495 Coco's Island site 23 marine clone S23_1361 EF5732
EU805181.1 gg_id:334084 It's all ranking aquatic 250 miles Panama City clone 6C233174 EU805
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EF573364.1 gg_id:232550 Coco's Island site 23 marine clone S23_1463 EF5733

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EU805034.1 gg_id:325083 It's all ranking aquatic 250 miles Panama City clone 6C233016 EU805
EU805307.1 gg_id:349966 It's all ranking aquatic 250 miles Panama City clone 6C233309 EU805
GU230237.1 gg_id:589643 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A GU230
L10935.1 gg_id:5908 marine clone FL11 L10935
EF573024.1 gg_id:227598 Coco's Island site 23 marine clone S23_1123 EF5730
EU802266.1 gg_id:305764 It's all ranking aquatic Northeast Colon Panama clone 4C229568 EU802
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EU800659.1 gg_id:353974 It's all ranking aquatic Delaware Bay NJ clone 2C228821 EU800
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EU804553.1 gg_id:317436 It's all ranking aquatic 250 miles Panama City clone 6C232500 EU804
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AB186970.1 gg_id:105858 Mariana trough hydrothermal vent water 0.2micro-m filterable fraction clone MT-NE AB186
GU061794.1 gg_id:586928 and function microbial mesoscale eddy perturbations South China Sea clone CEP-DC GU061
EU638843.1 gg_id:329172 Ecological Role Firmicutes Identified Thermophilic Microbial Fuel Cells thermophilic n EU638
GU061451.1 gg_id:562584 and function microbial mesoscale eddy perturbations South China Sea clone CE1-DC GU061
GU061695.1 gg_id:581190 and function microbial mesoscale eddy perturbations South China Sea clone CEP-5m GU061
EU802853.1 gg_id:324621 It's all ranking aquatic Northeast Colon Panama clone 4C230237 EU802
EU801758.1 gg_id:345184 It's all ranking aquatic Chesapeake Bay MD clone 3C003127 EU801
FJ744964.1 gg_id:558120 Transporter genes expressed coastal response dissolved organic carbon surface wate FJ7449
GQ351154.1 gg_id:554776 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Eastern GQ351
EU800851.1 gg_id:336634 It's all ranking aquatic Delaware Bay NJ clone 2C229079 EU800
FJ745165.1 gg_id:561678 Transporter genes expressed coastal response dissolved organic carbon surface wate FJ7451
EU805345.1 gg_id:342433 It's all ranking aquatic 250 miles Panama City clone 6C233349 EU805
EF573040.1 gg_id:234332 Coco's Island site 23 marine clone S23_1139 EF5730
DQ071171.1 gg_id:130137 marine surface water clone Surf1.8 DQ071
GQ348102.1 gg_id:590363 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
GQ348021.1 gg_id:590277 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
GU061790.1 gg_id:571620 and function microbial mesoscale eddy perturbations South China Sea clone CEP-DC GU061
GU230229.1 gg_id:575529 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A GU230
EU804396.1 gg_id:344788 It's all ranking aquatic 250 miles Panama City clone 6C232311 EU804
GU119384.1 gg_id:576863 Threatened Corals Provide Microbial Habitats reef water clone Reef_H22 GU119
EU803151.1 gg_id:308243 It's all ranking aquatic Northeast Colon Panama clone 4C230583 EU803
GU230235.1 gg_id:546752 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A GU230

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EU237291.1 gg_id:252418 Genes transcripts Eastern and Western Equatorial North Atlantic Ocean microbial as: EU237
GU119329.1 gg_id:537039 Threatened Corals Provide Microbial Habitats reef water clone Reef_N05 GU119
DQ071149.1 gg_id:136073 marine surface water clone Surf1.23 DQ071
EU804835.1 gg_id:328677 It's all ranking aquatic 250 miles Panama City clone 6C232800 EU8048
AJ400350.1 gg_id:71838 dominates and dimethylsulphoniopropionate uptake after algal bloom North Sea f cyt AJ4003
EF572090.1 gg_id:233599 Coco's Island site 23 marine clone S23_189 EF5720
EF572828.1 gg_id:228464 Coco's Island site 23 marine clone S23_927 EF5728
EU800463.1 gg_id:310712 It's all ranking aquatic Delaware Bay NJ clone 2C228574 EU8004
EU544717.1 gg_id:262536 free-living particle- aggregate- archaeal and Laptev Sea 0.22-1.0 um size fraction (fre EU5447
EF574586.1 gg_id:235137 site S25 near Coco's Island marine clone S25_930 EF5745
GU061508.1 gg_id:552249 and function microbial mesoscale eddy perturbations South China Sea clone CE2-5m GU061
EU802748.1 gg_id:340698 It's all ranking aquatic Northeast Colon Panama clone 4C230118 EU8027
GU061812.1 gg_id:592029 and function microbial mesoscale eddy perturbations South China Sea clone CEP-DC GU061
EU804885.1 gg_id:310744 It's all ranking aquatic 250 miles Panama City clone 6C232857 EU8048
EU800228.1 gg_id:347716 It's all ranking aquatic Delaware Bay NJ clone 2C228250 EU8002
EF572698.1 gg_id:233393 Coco's Island site 23 marine clone S23_797 EF5726
EU800767.1 gg_id:336527 It's all ranking aquatic Delaware Bay NJ clone 2C228961 EU8007
GU230210.1 gg_id:576779 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A GU230
EF573605.1 gg_id:230554 Coco's Island site 23 marine clone S23_1704 EF5736
EU805370.1 gg_id:355973 It's all ranking aquatic 250 miles Panama City clone 6C233377 EU8053
GQ348121.1 gg_id:574611 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ348
EU800573.1 gg_id:336168 It's all ranking aquatic Delaware Bay NJ clone 2C228719 EU8005
GQ347984.1 gg_id:535016 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
FJ826463.1 gg_id:552472 Succession during spring diatom bloom Sea filtered surface sea water contrasting nor FJ8264
EU805242.1 gg_id:334464 It's all ranking aquatic 250 miles Panama City clone 6C233239 EU8052
GU230241.1 gg_id:548888 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A GU230
GU061807.1 gg_id:554292 and function microbial mesoscale eddy perturbations South China Sea clone CEP-DC GU061
EF572202.1 gg_id:228993 Coco's Island site 23 marine clone S23_301 EF5722
EF571907.1 gg_id:233932 Coco's Island site 23 marine clone S23_6 EF5719
EU804502.1 gg_id:348475 It's all ranking aquatic 250 miles Panama City clone 6C232440 EU8045
EU800005.1 gg_id:322498 It's all ranking aquatic Newport Harbour RI clone 1C227685 EU8000
EF572463.1 gg_id:232144 Coco's Island site 23 marine clone S23_562 EF5724
EF572449.1 gg_id:232239 Coco's Island site 23 marine clone S23_548 EF5724
GU230236.1 gg_id:555992 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A GU230
GQ347181.1 gg_id:591181 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
EU804475.1 gg_id:348011 It's all ranking aquatic 250 miles Panama City clone 6C232409 EU8044
EF572912.1 gg_id:233011 Coco's Island site 23 marine clone S23_1011 EF5729
EU802138.1 gg_id:312524 It's all ranking aquatic Chesapeake Bay MD clone 3C003551 EU8021
EU394584.1 gg_id:343762 Coccolithophore bloom dynamics shape (NE-Atl) northern Bay Biscay pelagic particle EU3945
GU061702.1 gg_id:536611 and function microbial mesoscale eddy perturbations South China Sea clone CEP-5m GU061

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EF573095.1 gg_id:230831 Coco's Island site 23 marine clone S23_1194 EF573
EU800405.1 gg_id:347665 It's all ranking aquatic Delaware Bay NJ clone 2C228495 EU800
AB193898.1 gg_id:108822 deep-sea vent clone MT797-NB06 AB193
GQ347809.1 gg_id:581936 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
GQ346920.1 gg_id:546028 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ346
GQ350110.1 gg_id:557752 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ350
EU800275.1 gg_id:332963 It's all ranking aquatic Delaware Bay NJ clone 2C228315 EU800
EU802779.1 gg_id:317970 It's all ranking aquatic Northeast Colon Panama clone 4C230154 EU802
EF573478.1 gg_id:233480 Coco's Island site 23 marine clone S23_1577 EF5734
EU800496.1 gg_id:354542 It's all ranking aquatic Delaware Bay NJ clone 2C228617 EU800
EU805076.1 gg_id:303685 It's all ranking aquatic 250 miles Panama City clone 6C233059 EU805
GQ347932.1 gg_id:579167 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
DQ071040.1 gg_id:133968 marine surface water clone Chl1.22 DQ071
EU805407.1 gg_id:349492 It's all ranking aquatic 250 miles Panama City clone 6C233418 EU805
EU802293.1 gg_id:335921 It's all ranking aquatic Northeast Colon Panama clone 4C229602 EU802
EF572302.1 gg_id:231569 Coco's Island site 23 marine clone S23_401 EF5723
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GQ347999.1 gg_id:593858 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ347
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EU802938.1 gg_id:312639 It's all ranking aquatic Northeast Colon Panama clone 4C230334 EU802
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EU802246.1 gg_id:337290 It's all ranking aquatic Northeast Colon Panama clone 4C229547 EU802

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U75258.1 gg_id:5904 marine planktonic clone SAR241 U75258
GQ337209.1 gg_id:583381 Hydrography shapes biogeography deep Arctic Ocean water depth 400m clone CB1 GQ337
FJ545627.1 gg_id:571962 Diveristy Seawater and Sediment Northern North Yel Sea seawater clone W2-52 FJ5456
GU230232.1 gg_id:550223 Multilevel approach along Rio de la Plata-South Atlantic Ocean coastal water clone A GU230
EU802776.1 gg_id:339096 It's all ranking aquatic Northeast Colon Panama clone 4C230151 EU802
EU799116.1 gg_id:308445 It's all ranking aquatic Newport Harbour RI clone 1C226678 EU799
DQ009163.1 gg_id:126283 bacterioplankton clone SPOTSAPR01_5m40 DQ009
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EU804225.1 gg_id:318608 It's all ranking aquatic 250 miles Panama City clone 6C232125 EU804
GU061813.1 gg_id:539918 and function microbial mesoscale eddy perturbations South China Sea clone CEP-DC GU061
EU802664.1 gg_id:336767 It's all ranking aquatic Northeast Colon Panama clone 4C230022 EU802
EU802586.1 gg_id:317606 It's all ranking aquatic Northeast Colon Panama clone 4C229937 EU802
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EU804890.1 gg_id:324490 It's all ranking aquatic 250 miles Panama City clone 6C232862 EU804
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GU061792.1 gg_id:582928 and function microbial mesoscale eddy perturbations South China Sea clone CEP-DC GU061
EF572150.1 gg_id:229222 Coco's Island site 23 marine clone S23_249 EF5721
EU802828.1 gg_id:312388 It's all ranking aquatic Northeast Colon Panama clone 4C230207 EU802
EU803133.1 gg_id:349290 It's all ranking aquatic Northeast Colon Panama clone 4C230561 EU803
EU802309.1 gg_id:328726 It's all ranking aquatic Northeast Colon Panama clone 4C229620 EU802
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DQ396078.1 gg_id:155140 deep-sea octacoral clone ctg_NISAA12 DQ396
AF353223.1 gg_id:22763 marine clone Arctic95D-8 AF353
GU061818.1 gg_id:582665 and function microbial mesoscale eddy perturbations South China Sea clone CEP-DC GU061
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EU005952.1 gg_id:261438 Major activity inside and outside natural iron-fertilized phytoplankton bloom 3 micro EU005
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EU801227.1 gg_id:335684 It's all ranking aquatic Chesapeake Bay MD clone 3C002474 EU801
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EU010144.1 gg_id:245665 Biogeography Singapore seawaters seawater clone B55 EU010
EU802276.1 gg_id:354196 It's all ranking aquatic Northeast Colon Panama clone 4C229581 EU802
GQ351241.1 gg_id:585642 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Eastern GQ351
EU802321.1 gg_id:346262 It's all ranking aquatic Northeast Colon Panama clone 4C229633 EU802
EF572632.1 gg_id:227854 Coco's Island site 23 marine clone S23_731 EF5726
EU802028.1 gg_id:317489 It's all ranking aquatic Chesapeake Bay MD clone 3C003417 EU802
EU800327.1 gg_id:326805 It's all ranking aquatic Delaware Bay NJ clone 2C228380 EU800
U75256.1 gg_id:5902 marine planktonic clone SAR211 U7525
GQ346815.1 gg_id:551062 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Saanich GQ346
DQ071172.1 gg_id:128312 marine surface water clone Surf1.9 DQ071
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GQ337168.1 gg_id:544967 Hydrography shapes biogeography deep Arctic Ocean water depth 410m clone CB1 GQ337
EU804826.1 gg_id:307486 It's all ranking aquatic 250 miles Panama City clone 6C232791 EU804
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GQ351217.1 gg_id:593777 Metagenome versatile chemolithoautotroph expanding oceanic dead zones Eastern GQ351
GQ337299.1 gg_id:555674 Hydrography shapes biogeography deep Arctic Ocean water depth 252m clone NB2 GQ337
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EU802856.1 gg_id:316482 It's all ranking aquatic Northeast Colon Panama clone 4C230240 EU802
EU801709.1 gg_id:307941 It's all ranking aquatic Chesapeake Bay MD clone 3C003069 EU801
AB187007.1 gg_id:106132 Suiyo seamount hydrothermal vent water 0.2micro-m filterable fraction clone SSM-N AB187
EU804121.1 gg_id:323752 It's all ranking aquatic 250 miles Panama City clone 6C232013 EU804
GQ337208.1 gg_id:571518 Hydrography shapes biogeography deep Arctic Ocean water depth 400m clone CB1 GQ337
GQ337194.1 gg_id:557396 Hydrography shapes biogeography deep Arctic Ocean water depth 1000m clone CB GQ337
GQ337176.1 gg_id:568404 Hydrography shapes biogeography deep Arctic Ocean water depth 1000m clone CB GQ337
EU802580.1 gg_id:332667 It's all ranking aquatic Northeast Colon Panama clone 4C229928 EU802
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EU802269.1 gg_id:318647 It's all ranking aquatic Northeast Colon Panama clone 4C229571 EU802
EU801648.1 gg_id:348015 It's all ranking aquatic Chesapeake Bay MD clone 3C002994 EU801
GU061225.1 gg_id:587129 Evaluation culturing methods cultivate previously Yel Sea intertidal beach seawater GU061
EU803197.1 gg_id:355010 It's all ranking aquatic Northeast Colon Panama clone 4C230639 EU803
GU061335.1 gg_id:545379 and function microbial mesoscale eddy perturbations South China Sea clone CE1-5m GU061
EU802749.1 gg_id:323099 It's all ranking aquatic Northeast Colon Panama clone 4C230121 EU802
EF572257.1 gg_id:234083 Coco's Island site 23 marine clone S23_356 EF5722
EU802486.1 gg_id:346488 It's all ranking aquatic Northeast Colon Panama clone 4C229820 EU802
U75255.1 gg_id:5901 marine planktonic clone SAR203 U7525
GU061422.1 gg_id:560097 and function microbial mesoscale eddy perturbations South China Sea clone CE1-DC GU061

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EU802844.1 gg_id:308027 It's all ranking aquatic Northeast Colon Panama clone 4C230228 EU802
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EU804375.1 gg_id:308733 It's all ranking aquatic 250 miles Panama City clone 6C232287 EU804
EU801132.1 gg_id:356060 It's all ranking aquatic Delaware Bay NJ clone 2C229469 EU801
GQ337177.1 gg_id:580656 Hydrography shapes biogeography deep Arctic Ocean water depth 1000m clone CB GQ337
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GQ337245.1 gg_id:550273 Hydrography shapes biogeography deep Arctic Ocean water depth 1000m clone CB GQ337
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AF479300.1 gg_id:111655 Orientia tsutsugamushi str. TH1817 AF4793
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U12462.1 gg_id:5956 Rickettsia sibirica str. 246 U12462
L36221.1 gg_id:5932 Rickettsia typhi str. Wilmington L36221
AF060705.2 gg_id:5958 Rickettsia honei str. RB AF0607
AB021128.1 gg_id:5952 Rickettsia sp AB0211
U15162.1 gg_id:5968 Rickettsia canadensis str. McKeil U15162
EU036982.1 gg_id:345789 Rickettsia raoultii str. Elanda-23/95 EU0369
L36673.1 gg_id:5976 Rickettsia parkeri str. Maculatum 20 L36673
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L36213.1 gg_id:5948 Rickettsia japonica str. YM L36213
AF172942.1 gg_id:112194 Rickettsia hulinii str. HL-93 AF1729
DQ100164.1 gg_id:145296 Rickettsia monacensis str. IrR/Munich DQ100
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AJ628417.1 gg_id:112673 Wolbachia pipientis AJ6284
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L02884.1 gg_id:6105 Trichogramma deion symbiont L02884

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AF304445.1 gg_id:43030 Wolbachia sp. Okinawa-RS32	AF304
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AF432339.1 gg_id:31853 template extracted sediment oil-contaminated nutrient amended beach microcosms;	AF4323
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FN431786.1 gg_id:562499 Erythrobacter gaetbuli str. AMV3 AMV17	FN431
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FN668376.1 gg_id:519549 Erythromicrobium sp. str. BF8	FN668
GQ284449.1 gg_id:541199 Erythromicrobium ramosum str. THWCS10	GQ284
AB013355.1 gg_id:6348 Erythromicrobium ramosum str. DSM8510T	AB013
GQ472116.1 gg_id:563240 Microcystis aeruginosa freshwater lake clone C-71	GQ472
GU321359.1 gg_id:546220 Porphyrobacter sp. str. NMC22	GU321

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DQ001633.1 gg_id:158572 Microbial seeps asphalt seep clone E91-43 DQ001
FJ997597.1 gg_id:421125 Altererythrobacter sp. str. KYW147 FJ9975
GQ264058.1 gg_id:588310 Structure Simulated Level Waste Site simulated level waste site clone WC3_156 GQ264
DQ676427.1 gg_id:219175 Archaeal and plankton freshwater pond suboxic freshwater-pond sediment clone M DQ676
DQ811848.1 gg_id:162725 mangrove soil clone MSB-3A6 DQ811
EU440968.1 gg_id:312240 Lutibacterium anuloederans str. PR54-13 EU440
NR_025816.1 gg_id:536261 Porphyrobacter donghaensis str. SW-132; KCTC 12229 NR_02
AB033328.1 gg_id:6344 Porphyrobacter tepidarius str. DSM 10594 AB033
AF465839.1 gg_id:99433 Porphyrobacter tepidarius str. DSM 10594 AF465
AM691106.1 gg_id:221235 Erythrobacter longus str. EG6 AM691
EU770257.1 gg_id:346240 Porphyrobacter sp. str. A2 EU770
FJ516793.1 gg_id:539522 semiarid 'Tablas de Daimiel National Park' wetland (Central Spain) unravelled biofilm FJ5167
EU276570.1 gg_id:362799 -DGGE and Approaches Analyze Effect Agricultural Practices on Soil agricultural soil c EU276
GQ016328.1 gg_id:493339 Topographical and Temporal Human Skin Microbiome skin interdigital web space clc GQ016
DQ399262.2 gg_id:156567 Erythrobacter indica str. YPM1 DQ399
DQ448852.1 gg_id:164787 Novosphingobium nitrogenifigens str. Y88 DQ448
AF465836.1 gg_id:79367 Erythrobacter litoralis str. DSM 8509 AF465
EF424398.1 gg_id:205122 Erythrobacter sp. str. SM115 EF4243
DQ985038.1 gg_id:181578 Erythrobacter sp. str. JL1020 DQ985
FJ674974.1 gg_id:516226 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6749
AF407713.1 gg_id:60586 Great Artesian Basin clone B10 AF407
AM176884.1 gg_id:161637 Recovery mangrove sediment clone SZB32 AM176
GQ476825.2 gg_id:534168 Altererythrobacter sp. str. R83-1 GQ476
FJ672518.1 gg_id:531645 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6725
EU118985.1 gg_id:243523 Novosphingobium yangbajingensis str. YBSY-12-2 EU118
EF520398.1 gg_id:231088 structure lakes: what controls ? acid-impacted lake clone ADK-BTh02-25 EF5203
DQ520195.1 gg_id:160939 shal China sequences hypertrophic freshwater lake clone ML-9-79.2 DQ520
AY576736.1 gg_id:139435 Erythrobacter sp. str. 5IX/A01/140 AY576
DQ304436.1 gg_id:147665 Alterierythrobacter epoxidivorans str. JCS350 DQ304
DQ532296.1 gg_id:171201 Geographical Variation Microbial Burden and Spacecraft Assembly Kennedy Space C DQ532
AM936339.1 gg_id:547624 Dynamics during identified and SSU libraries pilot-scale bioremediation process hyd AM936
AY922042.1 gg_id:114156 farm soil clone AKYG623 AY922
AM746688.1 gg_id:256823 etiology rosy discoloration ancient wall paintings rosy-discoloured mural clone A10 AM746
EU328097.1 gg_id:278897 dynamics during bioremediation crude oil contaminated moderate saline soil clone Y EU328
EF157081.1 gg_id:256321 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF157
GQ048646.1 gg_id:445570 Topographical and Temporal Human Skin Microbiome skin volar forearm clone nbu4 GQ048
FJ675449.1 gg_id:518998 Synecology primary and secondary habitats Escherichia coli O157:H7 feedlot surface I FJ6754
AB012062.1 gg_id:6337 Erythrobacter sp. str. MBIC3019 D1215 AB012
EU104298.1 gg_id:345528 Evidence Comamonadaceae determinants settling performance activated sludge clo EU104
CU914890.1 gg_id:445733 and dynamics adapted microbiota during degradation Tunisian zarzatine enriched se CU914

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DQ396158.1 gg_id:151573 deep-sea octacoral clone ctg_NISA006 DQ396
EF157121.1 gg_id:257302 Microbial natural asphalts rancho la brea heavy oil seeps Rancho La Brea tar pits clon EF1571
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EU571151.1 gg_id:266620 eubacterium sp. str. 7-22 EU5711
DQ011529.1 gg_id:122133 Porphyrobacter dokdonensis str. DSW-74 DQ011
AF361178.2 gg_id:31470 Sphingomonas sp. Pee Shoal AF3611
DQ860069.1 gg_id:166645 Symbiotic aerobic anoxygenic phototrophic colonial tunicate clone L30 DQ860
AF407721.1 gg_id:56491 Great Artesian Basin clone B37 AF4077
DQ856560.1 gg_id:240257 intestinal microflora Chinese mitten crab (Eriocheir sinensis) clone C3X DQ856
EU784670.1 gg_id:341576 Sphingomonas sp. str. FG03 EU7846
CU466791.1 gg_id:328857 Evry wastewater treatment plant anoxic basin clone CU4667
EU131005.1 gg_id:251580 Sphingomonas sp. BAC151 EU1310
DQ860058.1 gg_id:164190 Symbiotic aerobic anoxygenic phototrophic colonial tunicate clone C1 DQ860
EF516627.1 gg_id:225426 grassland soil clone FCPS560 EF5166
DQ985034.1 gg_id:182499 Erythrobacter sp. str. JL990 DQ985
DQ898300.1 gg_id:162144 Sphingomonas sp. str. MTR-71 DQ898
EU360289.1 gg_id:263034 Pyrene-Degrading Consortium Deep-Sea Sediment West Pacific and its Key Member EU3602
AY792292.1 gg_id:123759 Humic Lake clone CrystalBog5F2 AY7922
DQ454121.1 gg_id:156628 Porphyrobacter sp. str. J3-AN66 DQ454
FN397674.1 gg_id:460604 Erythromicrobium sp. str. AMV10 FN3976
EU283461.1 gg_id:277643 Model Early Earth Harbors Deeply Divergent Divisions Anderson Lake clone A553 EU2834
EU327998.1 gg_id:277975 dynamics during bioremediation crude oil contaminated moderate saline soil clone B EU3279
GQ060166.1 gg_id:459373 Topographical and Temporal Human Skin Microbiome skin toe web space clone nbw GQ060
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CU920729.1 gg_id:533603 Towards definition core microorganisms involved digestion mesophilic anaerobic dig CU9207
FJ444668.1 gg_id:569696 Cotton Rhizosphere cotton rhizosphere clone 2y-17 FJ4446
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FM873313.1 gg_id:367391 occupant source house floor dust clone FD01E08 FM873
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AF468357.1 gg_id:79915 Arctic sea ice ARK10031 AF4683
AF468353.1 gg_id:79611 Arctic sea ice ARK10016 AF4683
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DQ453142.1 gg_id:157567 Erythrobacter sp. str. CNU001 DQ453

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k__Bacteria	k__Bacteria	p__Proteobacteria	c__Alphaproteobacteria

o__Sphingomonadales	f__Erythrobacteraceae
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g__unclassified
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g__Blastomonas
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g__Novosphingobium
g__Novosphingobium
g__Novosphingobium

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s__unclassified	61372
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s__Blastomonas natatoria	60232
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s__unclassified	60892
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s__unclassified	61275
s__Novosphingobium capsulatum	60349
s__Novosphingobium capsulatum	60573
s__Novosphingobium resinovorum	60586